

<110> Byrum, Joseph R.
 <120> NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH PLANTS
 <130> 38-21(15598)B
 <160> 36935
 <210> 1
 <211> 147
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1

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 atggatggtg gntttggaca tttggat 147

<210> 2
 <211> 378
 <212> DNA
 <213> Glycine max
 <400> 2

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 agcaaccgtg aactgctccc tctgcacttg ggtttttaa tccactgaa actgacaacg 180
 ccaacctcac tggcgccctcc ctctcaccca aatcgccacc tcgagtggcg ccttccagct 240
 aggaccctgc aactgcaggt gctgcaacct atggttgag aagtgcaggt gctgaagtcg 300
 ctggtcaaac cagcgctttc atgcacctt ggacgcgcca ataagagctg cgatatgcag 360
 tcctttgtcg ccagtcac 378

<210> 3
 <211> 423
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 3

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aaatggcgga gttgcagcgg tgatcacaga agatagtgat ctaatagcat atggctgtcc 300
agctgtaaga actcctccaa tactgtgata ttgcgcattg aggtttactg ctnnttttgat 360
atctcgattt atttacttgt tcaactattca gttcataga aagcatgcat tttgggatat 420
aat 423

<210> 4
<211> 462
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 4

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aaacagtaca gcagatgaaa atggtgggtc gcattaacct tcacaatttg gcagcacaga 180
aatcggacta ttttctcaaa cgaaccattt aatggaagca gaatgaatga ggatgcagtg 240
ttactggtct ggtcatggtt cacagagctg gagaaaagat ttacagagca ttacaactac 300
tggtccagca acctatcagt agctntctgt aactagcata gatgggaagc tattgaccaa 360
atgtaacaat gtatctagtc tggattctaa cagagggacc ttcatcccac acagtcatac 420
tcctgtattc ttagtaccac ggtacttttc tataatataa at 462

<210> 5
<211> 394
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 5

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tcaaataccta aaatttgagc tcctagggag caaaacaatg tgtgtctcct agagagggca 180
 tcagctacca catttgtttt tccctttttg tatttgataa catatggaaa ttgctctagg 240
 tactctaccc attttgcatg cctcttggtt aacttgcttt gccctctaata gtacttaagt 300
 gattgatgat cactatgaat gacaaattcc ttggaaacaa ggtgtcgcaa cctacccttc 360
 ngcgggaggg cgacgcgtga ctgcgggat gcgt 394

<210> 6
 <211> 464
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 6

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 agttacaaca agtgttacac atgcttctat ttatagacta ggtagcttcc ttgagaagct 180
 ttcttgagga aacttccttg agaaacttct ttgaaaaaac ttccttgaga aggtagagct 240
 tagctacaca caccatctc ataactaagc tcacctcctt gagaagtttc cataagaaga 300
 ttcttaaaga agctagagct tagctacaca tacctctcta atagctaagc tcacctcctt 360
 gagatgggaa gctagagctn tgctacacac ccnctatgat agctaagctc acccccatga 420
 caaaatacat ganaatacaa aaaagatccc tactacaaag acta 464

<210> 7
 <211> 373
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 7

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 tacaagttcg cttagcacac tgtttcgtct cactaagcgc accgcttcag tccatcagct 180
 aagcgagaaa ggcacgcgt aagccgaaat tctaataatgt gcgctaagcg gtccagaatt 240
 gcgctaagtg cagcagcacg aacaaggcca cctatttaag cttgaaatca gattttgtga 300

aggagagtttg ggctaggatt cagagctttg catgtctaga gattctagag agagaaaggt 360
ccaatttcag aga 373

<210> 8
<211> 462
<212> DNA
<213> Glycine max

<400> 8

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tacctgtcgc aagggtttgt ggtttgtgct cctctgctga ccaccataca gacctttgcc 120
cttccatgca gcaacctgga gcaattgagc agcctgaagc ttatgctgca aatatttaca 180
atagacctcc tcaacctcag cagcaaaatc aaccacagta gagcaattat gacctttcca 240
gcaacagata caacctgga tggaggaatc accctaacct cagatgggtcc agccctcagc 300
aacaacaaca gcagcctgct ccttccttcc aaaatgctgc tggcccaagc agaccataca 360
ttcctccacc aatccaacaa cagcaacaac cccagaaaca gccaacagct gaggcccttc 420
cacaaccttc cctcgaagaa cttgtgaggc aaatgactat gc 462

<210> 9
<211> 421
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 9

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atcttcttta atctctttca acattttcaa cagatctttc tgattttattt cctttcatct 180
ttctaaaagt ttttgttcaa tagtttctct tccaagaaaa gttctttgtt caaaaacttc 240
agctattcat ctttttcatt ctcttctccc tttgccaaaa gaaggaagga ctaaccgcct 300
gaattttttt gtgtctctct tctcccttac aaaagattca naggactaac cgctgatat 360
atcttttgtt tccccataca aagattttaa ggactaactg cctgagaatt ctttgtccca 420
a 421

<210> 10
 <211> 404
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 10

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 tgcagaggat tgagctccat atgcatgcat atatgcaaca tgtggccgac caataggcgg 180
 ccaatcatag gggatagggtg cagctgaatc agagctttta ccagtacacc ctacgtcagc 240
 agagctagga tcccagccct tactcgtggc ttactccga gtagtttggg gccacagttg 300
 catggcctgg agataggecc aattttcaag tagggacaag accctcaaag gccccaggag 360
 ttgaagatgg agctcaagaa gacgacgaca tangcgatgt gatg 404

<210> 11
 <211> 284
 <212> DNA
 <213> Glycine max

<400> 11

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 attaacaact ttcgtttgcc catctgcttg tgggagacat gtggctgaaa ataacaattt 120
 agtgcccaac ttgaccacac caggactacg caaatggctt atgaacttac agtccttacc 180
 actaacaatg ctgcttggtg aacctatgat gtcacaatc tccttgagga acaaattagc 240
 cacatgggaa gcatcatcta cttctttaca tggaataaaa tgag 284

<210> 12
 <211> 373
 <212> DNA
 <213> Glycine max

<400> 12

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 actaatatct taagtcctat atataaatct tctctttgag cacttcttta tagctaattg 120
 gaagtacatt ataaccacca atttgataag atattgagac ataggcaaac cactaatcca 180

atgtacatac tgcagtgcac acatgttgta tccggaaagg attcacatgc atagagacat 240
 tgtgaaccca agattcctac tatgtttgtg gcaatggaaa gagttaacaa acagtgttga 300
 aaccaccctt tggttaatgcc tatgaagaca aacttactgt cacacctata ctaaaacacc 360
 cctaacatat act 373

<210> 13
 <211> 339
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 13

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 aagattgtc ttgcctttga ctttctctat ctctctcgn gatttttttt atttgagcaa 180
 ccgttgatta tccggtaggg gtggaacttc gtatatgtct ttaatatctt cccatagatc 240
 acaagcatca agatagggtt ccgttctaata agcctagagg tggtaatgtt ntccattgaa 300
 tagtgaaagc ctatgaagca cggacaccct agtccctta 339

<210> 14
 <211> 397
 <212> DNA
 <213> Glycine max
 <400> 14

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 gataattcga gatcacttga aattagtga aaaaattgtt tccgtgaaga aaattcaagc 120
 caaggcggtt ccgtgggtga tttcggaag attttcaacc gttcttcgac gttcttcggt 180
 cggttcttctg cggttcttcag ttttcaaccg gtaagttccc gaaatcgaac ttttcaattc 240
 attctatgta cccttagtgg tcttcatttg ttttcacgcy cttttatttt cgtttcattt 300
 actttccgta cccctttttg acgtgctcta gtcatttact taagtcatgt tctcgcccta 360
 tcaaaaaata aaataaatat ccactgatca tttgagt 397

<210> 15
 <211> 384

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 15

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gagttgggga ggattgattn tggaacttgt cgtgggtgcag aagttagttc aagtgcgaac 120
actactagaa aaagagcttt ttgcgatgca cttacgacat cggccaaca aaactgtcga 180
agtatattaa atgggtgcatt tgtgtaatta caacgaaagt gtgcaccttg ccaattttat 240
ggttgacatt ggcacaactc ccttgaaggt tgttggaagg gactcgagag tgaggaaaact 300
agacagcttg gggatttctc gatttccgta acatacttaa tgctctcaca acatagtgga 360
gttagggtag taaatttcac catt 384

<210> 16
<211> 243
<212> DNA
<213> Glycine max

<400> 16
cgaaccgcac cctactttat acggcgacaa acatgtggat atagacaaac atgcgctgac 60
ccgtctcagt gtcatgccta aggctagctc agcatgagtc caactttagc tagcgcgatt 120
cataatgagt tgtgccacat tttgcctata agtaggtgag gcgatttttt tcaaccaatt 180
agactcta atccatgggtgga tcaagttgac tcacaataat aataataatc tttttactta 240
cct 243

<210> 17
<211> 331
<212> DNA
<213> Glycine max

<400> 17
agctatatat cttttcttct tggttctgct tgctctgtag tgctttgggt ctatgctatc 60
cttttatatt tcatactatc tttgacacat gggactaaac attgaacagg tgggaagggtg 120
gtccagccaa aattccatgg tccttttaaag aagaagagac tatgtatctg ctggagggaa 180
aagtgagggt tactgttgaa gggctctgtt ggtcttttga aattgggggt ggtgatttag 240

ttgtcttccc aaaaggaatg aacattactt gggaagtgat tgaaactgtg aagaagcact 300
acagcttgaa aaaataatga tgtgtactta t 331

<210> 18
<211> 443
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 18

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gagaaatggt agcaatatac taccgtatga cactgcatca cacactttat tatttgccac 120
aatttattgg aaatcacaaa attttgtggg ttctgttact tatttaatga acttcactcg 180
tgattttgga atttctaata aattttaacc aataataata ataataagagt gtgttactta 240
gaagggcatt gtattgctag cactcctctt gaagtatagc atacaaacat gaaaggaatt 300
ccattttaag tattatcctg taccanaacc tcactttagt cccaatttt ggaaatcaca 360
gttcttttca ctgacaaatg acttacagtt ntagttaaaa atagggatta acaagagtgg 420
agcatacaag accaggaggg act 443

<210> 19
<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 19

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tcttagtatt tctttcaagt attgtgcta ctctagaaac aagaataaga gaaaattaaa 120
gggaatgcag ttagtcacat gatataattga gatttgtttt ttttttcttt ttttgtaatt 180
gataaaaaag acaaaattgg tgtcttttgc attaaggggc ctttcagaag aacttgtgaa 240
aataaattga tcagtttaat ttccttatac ttcaagtga aagattttta tactatgaac 300
taacaaaaaa tcctcctatg attnttaata taattattat aaaattacca tacatcataa 360
tttgagaatg tagaanacat aaacaacgtt tacact 396

<210> 20

<211> 459
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 20

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 acccttaaaa caaaaatggc atacaacctc ctccaataaa tacaacatc aatgtaaatt 120
 tagagcaagc ttatgcgcat attttcttac gaacattcac tcgcacaaga tattcttcta 180
 actaagaaaa atgcacccat gcacaatcaa ggcactttcg ttacctacat tatttgtatg 240
 tacttccaag gtgtactacc tacaccacat gcatttcctt ggctaaattt acatacatgc 300
 atgctcaaag cctcttggt accaaaagtt gcacacatgc aaactttatg atgaatcttg 360
 gctatctaca caataagggtg ctacacttca tgctttatat caagtgtttt actaccagaa 420
 gccgcatgcg aatgtcagta tattttcttt tgccgacta 459

<210> 21
 <211> 328
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 21

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 cttaggcact tctctctctt tcgaatttgc ttaggaaaat tgtttccgtg aagaaaatcc 120
 aagccgaggc gttccgtaa cgtttccgtg agtgattttg cgaaggtttt cgaccgttct 180
 tcgacgntct tcattcgttc ttcacgntc ttcagtcttc aacgggtaag tacctcatc 240
 caagcttttc aattcattct atatacccg nnggggccac attatggttc atgtattatt 300
 attctcgntt catttactct ttataccc 328

<210> 22
 <211> 391
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 22

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taagagtaac gtcccactgg taaaactaac tttccaaatg tttgccttcg caggaatggc 120
cccaggaag cttgcctcaa agaggteccag gaaggacaag gcggccgaag gaactagttc 180
cgccccggag tacgacagtc accgcttttag gagcgttgta caccagcagc gcttcgaagc 240
catcaaggga tggtcgtttc tccgggagcg acgcgtccag ctcatggacg acgagtatac 300
tgatttccag gaggaaatag ggcgcggcg gtgggcacca ttggttactc ccatggccaa 360
gtttgatcca gaaatagtcc ttgagtttta t 391

<210> 23
<211> 352
<212> DNA
<213> Glycine max
<400> 23

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accttatgaa aaagtatgga agatagtaga agagtgtaga gactcctaga atgtgtggag 120
tattctagag aattagtctc catcctagga tacaagtaat ctccactatt tattgtggag 180
gtggagtagt ataaataaag gtaggatacct tcattcctaa aaaatctaag tagagagtct 240
ctctgagaga gaagataaat agcttttgaa gtctctatcc tcaaacataa gtaagcctct 300
ctgagagaga agataaatag cttgggaagt ctctatcctc aagcttgagt ga 352

<210> 24
<211> 438
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 24

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tattcccatt ctccccgctc aaccttgaa tcgaattatc tccagccacc caactacacc 120
caaaatagag gttcagaaag gaagcaaact aacactgcc aataaccccaa gttccagggt 180
ttaggtggga attatactat tatcaaaacg ggttctacaa cctcttatag aagcataact 240
cttgcaaaca cttttaatag taaaaaagaa aaaaaaact ttacgtcaca gaactcacta 300
ccaagtgaag aaccaccaac attgtcggtt tgtactctgc agtctgcacg tgtttccata 360

aacagaaaca cattgatttt aaattaatta atcgattaat actaccatca agtagtacca 420
cccctatatt ctttctta 438

<210> 25
<211> 104
<212> DNA
<213> Glycine max

<400> 25
ccacattatt tccatgacac aaattgcaaa atgatgattt ggaaacttca tgcaaaactg 60
gtcatgcatg cacctatgca gacactcaag tgtcaaattt ttat 104

<210> 26
<211> 386
<212> DNA
<213> Glycine max

<400> 26
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aagagttagg tctagccacg gcccacgagc atagaatcac ggatgagtat gctcaagtgt 120
atgcggaaaa agaggctaga ggaaggggtga tcgactcttt acaccaagag gcaaccatgt 180
ggatggatcg gtttgctctt accttgaacg ggagtcaaga acttcgccga ttgttagcca 240
aggccaaggc gatggcagac acctactccg cccccgaaga gattcatggg cttctcggct 300
attgtcagca tatgatagac ttaatggccc acataattag aaatcgttag gaaaattgta 360
tggtctctca gaccttgact ggatac 386

<210> 27
<211> 379
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 27

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cttaagcact tctctctctt tcgaatttgc ttggaaaaat tgtttccgtg aagaaaatcc 120
aagccgaggc gcttccgaaa cgtttccgta acgtttccgt gaggaatttc gcgaagggtt 180
cgaccgttct tcgacgttct tcattcgttc ttcacgttc ttcgatcttc aacgggtaag 240

tacctcgaac caagcttttc gattcattct atgtacctgt ggtgggccac attgtgggtc 300
 gtggattttt attctcgntt catttacttt ctatacccc ttttgacgtg gcttaagcca 360
 tttatttaag tcatttctc 379

<210> 28
 <211> 395
 <212> DNA
 <213> Glycine max

<400> 28

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 cctagtctgt ttgcgaagtt tgttcaggaa catgggatta ttgccagta cactatgctt 180
 ggttctccgg atcagaatga tgtggcagaa agaagaaacc gaactttaat ggacatgata 240
 aaaagtatga gaagtaataa aaaacttcct caattcttgt ggattgaagc attaaagacg 300
 gttgtatata tattaaccg ggttccaacc aaggctgtct taaagacacc tttctagtta 360
 ttcaaagggt ggaaaccgag tttgcgacat atatg 395

<210> 29
 <211> 408
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 29

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 cgtcagggcc tctccctcct gattcaggtc caaccanaa aacattntag cacacagact 120
 ntatctatga actgtacaaa atacacgact cctcaattgt tctcaaaata attttatcta 180
 atcgcgcttg tgattaaact cgtcagggtc caacagtggg tcccatcata atactcgcca 240
 cgcattaact cgtcgccctt agattcatag ttcacaaatc agggcacaca acatctcaat 300
 gcacatatat attacaagtc aatacatact caatttatca catacatttg gtctcaatca 360
 cagtgggtata atctcaattt aacatgttat cacacctcat gaatcata 408

<210> 30

<211> 443
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 30

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 agtctgatat ctcaagatta catttgaata aaatcatgag gttgaattaa ttgaaatact 120
 ttaagggttat agcaaaaaag gtttcagcta aaacaaatgc aaggcagcgt aagaaataaa 180
 ttactacatt agcaataacg cttaaataatc tacataaaca gaatattcca taaagattat 240
 atttaagccc catgctgaga tgcaagtaat atgctgtttc atatttatca aaatatagga 300
 atggaaatga tgcaggaggc ccacagaatt aagtcataaa cctgaactca actacatctg 360
 tgcatacaca taaaccanat cctaccattn taattntaca cctcccccacn acccacaatg 420
 aatatggcct aaggaatata tcg 443

<210> 31
 <211> 398
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 31

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 tgcttgaaag agttccttgt caggactttg tccttaggat caccaatgat ctgggactct 180
 agatgatctt tttgtagcaa tcatccagtt ggttctctga cttatagagg ttgatcatcc 240
 actggtgagt tggacgcata ttggtcttga ctggacctag caacatattt cacgatattn 300
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 tcattaagtc ttacgtgaat ggctcttcc acagtcac 398

<210> 32
 <211> 458
 <212> DNA
 <213> Glycine max
 <400> 32

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 catcaattcg gctttgacta ggaacacccat cattccctct tctcctcctt tcttcttcat 180
 tatgatctct attctccatt tgatccaacc tctcatggag cacatcatct cattgtttca 240
 ttaacctctc caaatgatgc atcacagctt gcatttggaa ttgcgaaagc cccactccat 300
 cattaggatt tggctctgcc atctcataca aacacatcag acgtatcaag acaattatag 360
 ttgctgtttg aatacctcac tcaactcaagt gtatcacaca attatggttt ttctctaagt 420
 aaacactctt gccttctacc actctaattc cacttgag 458

<210> 33
 <211> 382
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 33

agctttgata taatgctttc tttgtattta taggtataga ggatctagcc acttgagagt 60
 gtcttgcccc gactcagtac aatgcattat attagcgcga catagctttg gtctacattg 120
 aattagtata tatagctaca ccttattcaa tccaagaatg gaaaaaggga agcagtgggt 180
 aagaatgatc atgggtcaag cttgagtcaa gtaatctgat accaaaccga atgaaaatgg 240
 ttaggttgag tgtgttaa atgtgactgtct caaccacaac taatcacaag tgggatgtgc 300
 ctaagtcacg tcattatatt aaagattctg tgaatgaagg aagaaaaaac acacanaaat 360
 aggggtagag tagggaaaag gt 382

<210> 34
 <211> 460
 <212> DNA
 <213> Glycine max
 <400> 34

cctgagtga acaatgagac tcttcacagt taaatttgaa tttcgacgtt catggacact 60
 ggtaatcgat taccaaaaaca ttggaatcga ttatagcctt ttgaatatat tgggaacgtt 120
 gtaaatcag tttgaaaact ttttcaaact cttttagcta ctgagaatcg attacaacaa 180
 tatgaggatc gattaccaga gagtaaaagc tctttggtaa agattttgtc aaaaactcac 240

gagctataca acgttgagaa aaaacctttt taatacttat attgatagag tgtttgatac 300
 attctcaaat gttgaatggt gaatcttgat cttgattctt gagaactcga gtattgagtc 360
 ttgattatta accttgatgc ttgatgattg acatcatgaa tcgtgaatct tgatacttat 420
 ctgaaggctt tcttcttgag tcttgaattc ttgattcttg 460

<210> 35
 <211> 320
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 35

agcttgtaga gcttgagttc taagaatgag ctgagttatt ttgaacccat tntgctgttt 60
 atttttccta aaatggatta tgaatatagc ttggatacta tggaaacaaa aaatgagtat 120
 tctgatgaag gtgaagtagt accgggttta tccattaatc cggttagttt ttgcgcaaac 180
 attgttccaa tgattaatgg taaatatggt ttgcattacc aaagtgaatg caattctaca 240
 aagaattgca gtcctattgg ggggatgcc accaattgcc ttatattgtg tctttgaaag 300
 aactggtttg ttagatgtaa 320

<210> 36
 <211> 368
 <212> DNA
 <213> Glycine max
 <400> 36

tcacattcac tatcctctac atcatattca aacttgacca aataaatagt acagtcattc 60
 cgactcaaag aaggacatct aagtctcata caattaatat agaacctata tcctaattgtc 120
 acatcctatc aaagcgtggc gctaccgctt cctctagctt gaggtcttct atagtcattc 180
 acctattcat ctgctacccc gaacacagag cttgagatca tcacaggatg cgaacacaaa 240
 cagcacaccg ggagtggatg atcacacttt taactactat agagaaacaa cacaacatat 300
 atgagccgaa gacgatttac ttaccatata tcacattatt tcatgacttt gtccttcac 360
 gatcacac 368

<210> 37

<211> 414
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 37

agcttatcat ctatttatac tacacatagc aatgaatact tattctggat cactgcttaa 60
 tgaaattaag gatctgttta ttttacattg taaacaggca aatgtattaa ttttaagattt 120
 gtatctaaat tgttctcaac tatatataat tatatacatt ngtattaaca tgtataattc 180
 tagcatacag gcagttttta gaaatatacc atgtgactcg atcatgtgta tgtttgagtt 240
 gataatcttc ttggaggagt gttaagattc aacacatttg tattaatatg atgtgagagt 300
 cacaaattnt tatcattttc aatatatttc aatcaataac aaagaatata tttaaagaga 360
 ctcaacaaat atgtctctac tgtttctcac attattaata taatcaattg cttg 414

<210> 38
 <211> 288
 <212> DNA
 <213> Glycine max
 <400> 38

gttccaaaga ggtcttcggc attacattca aactcgatcc attgtcgata agtacatttg 60
 cgaccacgtg tgccgtacat atcaccgacg catgtacagc cttgatgtgc cctctcctct 120
 caacgggaat aacttcttgc acgaacgcga tataattgcc gatggctata tgattggcta 180
 tgccgatcag aacgatgcgt gagatataat gagctacact gtcatggaca tgatccataa 240
 tctgagacgc actgatacac tccctcaatt cttgtggatt gaagcatt 288

<210> 39
 <211> 317
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 39

agcttgtcac tgtggaatta accaagtctc tgagggacgc gttcaaggat aattcgaaat 60
 catatgccat agcgatctct tcaactctgac catcaataat tactaccatg gtttctatct 120
 ccaagatgtc aaccagaact gctacctttt gcctgagctt gaaggacgta tatgacactg 180

aatacaccaa taagtaatta atactacaat ctaatagctt aatacttaac agttgacaca 240
aatattaatt ntcattacct taatgttctg aagtgcacgt ctaaagtttt gtgcataaca 300
gtatcgccac taaaacc 317

<210> 40
<211> 435
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 40

cactatctat acttagcttg caatctatgt gcaccagata gctaataagac ttttgtattc 60
ttncgttatg agtatgatat ctcaagatta catttgattc acctcgtgat gctgcataaa 120
ttgagtatac ttcacgcacg ttgcaaatat aggttttagct taaactgatg cgctgcaccg 180
caggaaatag agtactacat tagcagtaac gcttaaatat ctacattatc agaatatccc 240
ataaagatta tattttaatcc ccacgtgat atgccagtaa tatgcctgct catattgatc 300
acaatatatg aagggatatg atgcacgatg cccacagaat taaatcatag acctgaactc 360
aactacatct gtgcatacac ataaaccaa tctaccatt gtaattttac acctcccca 420
aaccacaat gaata 435

<210> 41
<211> 285
<212> DNA
<213> Glycine max
<400> 41

agctatatat aagctcttct ttcacagat gctccgaaaa tttaacttct tgttgtgata 60
attagggggg agcagtttat aactggattt gtatctgaca gagagaaatc ttaacacaag 120
tcactctgac actcttattg tataacaaat taaggccact gagttgagtc cagctatcca 180
aaagctgtag gaataaaaaa tctattaaga gcaaacacac acctcgacct gtgttatgca 240
agtaaaggt aatgtaagcg actacgcagt agtacgggtt ggtag 285

<210> 42
<211> 452
<212> DNA
<213> Glycine max

<223> unsure at all n locations
 <400> 42

ttgatgatag anattaattg aatgatttat tattaattct tcttaatggt tggacaatat 60
 ttccaatatt agaaattaca aactcaaaag aaaccaataa ataaattcct agtaacaaaa 120
 cttgtttttt ataatttcat atttatcaat atttattata taaataataa taaattataa 180
 ttaaaaaatg aataagtatt atggttagata tttttataat aatataagat aatatctaata 240
 attaaaaaat atctatcaat gagatcggtc acttgtgtta gctaacttac atgaaaagtc 300
 aatgagatct gttacttgtg ttgcattggt gtagacgaaa cttgaacatc attagcaatt 360
 atcaagggtc tctatcatc acataaagta tgggtttgat acttaacaat aagcagacca 420
 tcacagaaag gatatgatag cactctgact at 452

<210> 43
 <211> 335
 <212> DNA
 <213> Glycine max

<400> 43
 agcttctact tatgtggcag ggcgggcttc cttcaccttc ttgtctccaa cgcgaaacttt 60
 gaccattatt cttccttccc gcgatgcttc ttttcatgtc cgcttgagtg ggcttatagc 120
 ctaaaccata cttcccacga tttccttggg tatttatcag gctagtatatg ccgcccgtgt 180
 tttttcctaa acccatcccg ggttcataac cgttcccaa cataactcgg gccatcatta 240
 tcgctgcac ggacagacaa ggcttgccaa agagggagtc cacggaggaa atgctgacca 300
 cctcaaaaga ctggaaagca gtttctaacg attct 335

<210> 44
 <211> 434
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 44

tgcctcanag agatccaaga aggataaagc agctgaagga actagttccg ctctgaata 60
 tgacagccat cgtttttagga gtgctgagca ccagcagcgc ttcgaggcca ttaagggatg 120
 gtcatttctc cgggagcgcac gcgtccagat cagggacgac gagtataccg acttccagga 180

ggagatagtt cgccggcggt gggcatcgct ggttaccccc atggccaagt tcgacccaga 240
catagtcctt gagttttatg ccaatgcttg gcctacagtg gaggggtgat gagatatgcg 300
atcctgggtg aggggggttag tggatcccat tcgatgcgga tgctctcagc cagttcttgg 360
gatatccttt agtgctggag gagggccagg agtgcaagta tggccaaagg aggaacccgg 420
ccgatggggt tgat 434

<210> 45
<211> 408
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 45

agctntntgg agtagaaaca tgggaccaac tcattttatt tcaaaaagga agtcatatct 60
agtcaaggtc tgagagacca tacaagtttc ctaacgattt ctaattatgt gggccattaa 120
gtctatcata tgctgacaat agccgagaag cccatgaatc tcttcggtgg cggagtaagt 180
gtctgccatc gctttggcct tggctaacaa tcggtgaagt tcttgactcc cattcaagggt 240
aagagcaaac cgatccatcc acatgggtgc ctcttggtgt acagagtcga tcaccctctc 300
tctagcctct ttttccgctg atacttgagc atagtcgccc gcaatcctat gctcgtgggc 360
cgaggctaga cctaactctt tcttgcatg atagctagca tgggtggct 408

<210> 46
<211> 74
<212> DNA
<213> Glycine max

<400> 46

tctcaaggaa gttttctcaa gagagcttct caaggaagct acctagtcta taaatagaag 60
catgtgtaac actt 74

<210> 47
<211> 358
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 47

agcttgaagc atctcatcat aggagatgat gatcacaaag aagcanagaa gggtagtagt 60
aataataata gtaataataa taatagtgat catgagagag gtggcaaagg aggatcaagg 120
aactcacttg gtgaacactt cacagaggaa gagaagcagc ataatcttca gctgggttagg 180
atgcaacaga ataaggacaa cctccaaggc ttgaagttga agaagttggg gcgctggttac 240
gccaaagttt tggngcattt gatgaangct aagcgtgatc ctcatctang tgggtgatgct 300
gggaaaaaac ctgtnttcaa gttatcagcc tagccaggga aaattttgga gttttact 358

<210> 48
<211> 407
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 48

tgcatttcac gagcatcact cttntctctt cctcttgtg ttgtgacatt ntcctattga 60
gaatggaaaa ttatttccat accttcaaat ttgcaaagaa atgtatccca aaaatgctac 120
tctaaaatag gaagagagtt tgctcttttg ttttcgctg gtccattaat atactaaatt 180
agaagtcatt aaacaagctt tttcaactct cataattgtg gcatttacta ttgaaggtag 240
gggaatgatc ttaaacggat tgaaaatatt aagcaaggaa aattactgca gaaattctta 300
acaaatgaaa tcagaagtca ctttctctag acgctgaaga agagcgggtn tgaattggcg 360
aacaccacgc ccacttgatt gcggatctag tctgtgagct agttcaa 407

<210> 49
<211> 420
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 49

agctntaaga cactagtctt cggccaactn tgttttaaca ccaaggcaat taccatgggtt 60
gctcataaat ccatatttac agaacaaaat tngggcatgg ngctcaatac aaatcanaaa 120
gaggttctaa atatgtatta gactaacaac ggcattccat tagacaaaga gagacttagt 180
tctctaagaa tcaaattcgc atgcaaattg aaaattatag gatttggaaa atcatcacct 240
tttccacact atctttactc ttcaaaaccg aanatgattc caactcttct cttttcctta 300

gagagaaata catgaagaaa ggatggatga agattattcc tgcacccaaa cggagattct 360
aggagcttan naattcactc tttatnatat canaatacaa ggaatcttan aaattactac 420

<210> 50
<211> 423
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 50

tcattgtttaa tcatacattc aatcttatta tagcaganaa ctcaaataaa aattctatta 60
aaaggtagca ctctgtggcat gagtccttat aaatatctac tacaaaagtt ttaaatactc 120
tgtccatgag gaaagcagtc cagtatcttc caatactcta tccattgatc catatttgac 180
cttttcccat accctccata tccaatgcc aatgggtcattc tccttcggga gcatcaaaat 240
aagtctgcgc agtttacgga ttgacttcat cagttntcaa cctagtggta tttttttcag 300
tctaagttag gacattacta tgcaagatga ccttggtcca tgtcaatggt tgagttcttt 360
gaacaactat tgctgactgc accacgcaac actatagata tcatttggag aggcaacatt 420
cat 423

<210> 51
<211> 401
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 51

agcttgcttc tacaatagtt tctctttacc tcattctgaag agttacgaag gttccataga 60
agaagaaagt ggctccaatt gcagaagaac cataaggctc tcggttacat tattcattag 120
tcttgctcgt gattgattga tgtaaagatc ttgcaataat cgtcgaanaa caagtagatt 180
aggagccata tacggattaa ggtatttcat ctaatcttta ataatgaggc atgttgtaaa 240
tcctagggct tttggtagat tgttctaggt tacgcacatg ttgaattnta gcttccgcat 300
aaagaataaa gaatacggat canaagttaa aattctaaca actataagat gaacatcagt 360
tgccaaattc tttgcacatg cattggatnt aaagaacaaa t 401

<210> 52
 <211> 464
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 52

tagactaagt tcagcctacc atcctcagac tgatggccaa actgaacgga ccatttagtc 60
 attggaggac cttttgaagg cgtgtgtctt agagcaaaaa ggaagttggg agagttttct 120
 tccattaata gagttcactt ataacaacaa ttttcactct acgattcata tggctcctta 180
 tgaagctttg tatggtagaa ggtgtaggac acccctatgt tggttaaagc ccggagaagg 240
 ccttacctta ggaccggaag tgggtacaaca aaccaccgag aaagtcaagt taatccagga 300
 aaggatgagg actgctcaga gtangcaaaa aagttatcat gataagagga ggaaagatct 360
 gaaatttgag gttggtgatc atgtattctt gagaatcact ccgtggactg gggttggtcg 420
 agcattgaaa tcccgaagc tcacacctca ctttatcgat cctt 464

<210> 53
 <211> 342
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 53

agcttttagga taactttata tttgnngaag gacagtagac ccacaattgt atttcatcca 60
 cagaactcta acccacaatt gtatttcacac cacagaactc taaccacaaa catactagga 120
 tcttcaatca tatatcatcc caacatgaat aaataagact gattaagagt cctcaagctc 180
 ttaaatacaa gcccccttg atctttggat ttacaaatca tctaagaaat aagatgaggt 240
 gttctgttat ttgcatcact tcctgagata aagtctctgc agaagctttc aatctcatta 300
 cgaatagcaa tcgtaataaa ggttgcttca agaacataag ta 342

<210> 54
 <211> 460
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 54

ntntattgat tnnTTTTatg tcttgaccaa gaattagaga ctaattcttt gaaatattga 60
 gatttattta aaagataata tctcccaaaa aaatgttatt atattcaaac tgattaatat 120
 gaacatgaaa aataaacaaa acgccttata aaatttagat ttaggaaatt atttttaatt 180
 actttaggaa atgttattga tgtcacattc aaaaagtatg tgaaaggatg agtgataaat 240
 cataaaattt ggctgctata agttatatcg ataagattaa atttaatttt taactcaaga 300
 attaaggaaa gctttcataa aaagagaaaa atcaaatttt catttgacat gataatgggt 360
 agagcctaaa aataaaatat aaattaaaaa tatacatatc aaaatacatc taaattaatt 420
 aaataaaaag tactaaattc attggaaact agaaaatgga 460

<210> 55
 <211> 371
 <212> DNA
 <213> Glycine max

<400> 55
 agcttgta caacaacacc aacaaagtcc aagaatccct ccataacaat gatgctcaat 60
 accaacacgc tctttccac cttctctctg tctctcaggt atatttgcaa ttcattgcata 120
 ttgatatgct catatgcaaa aactagtctt aaattttatt cttgcgtatg gtgtttgttt 180
 attatatgca tagtttgta atcttcctta aaactttatt ttaatattaa tggatgtat 240
 tgaatgtttt taatggttga gataggtagc actgacacag aagtgtgaa tttattggca 300
 gttgaaagga gaagagatac ttgagcaatt cgaagcttct agttcttctg agccggtcgc 360
 ttctataact c 371

<210> 56
 <211> 459
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 56

ntgaaccatc cgtcccatag gagcccaact ntgcaggatg aaggggtcaa acaaaatttt 60
 gatgaagcta caaaggagaa ctttggtata ggaatgggct taattgtaga aaccatcacg 120
 gggagatcct aggggttgcc atgagcatgg tgaaccattg ttatcgaaca cacatggcat 180
 aagctgtagc tttcaaattg gccttaaaga tagctaagga cctctttttt tttttttttt 240

ttgacattgt	catggaaaca	gattgcttga	agattgttta	gacttggcac	aacacaagga	300
agttttcaac	ttcctatttt	gaaggcatcc	ttgatgattg	tagagagctg	cagagtagag	360
gttttcatac	gttcaaaatg	tcttttgtaa	agcatacagg	aaacaaagta	tatgggttcgt	420
tagtgaatth	agctcttggt	tttagggaac	gttattgga			459

<210>	57
<211>	423
<212>	DNA
<213>	Glycine max

<211> 247
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 61

 agctttgtga atctcctgat ctacgcccac ccgatgatac tcaaccattc gaagntgaat 60
 gtgatgctag tggaattggc attggagctg tcttgataca caacataatg cctatagctt 120
 atttctcgga gaaagtggga agagccttgc tgaattattg cacctatgac atagagatct 180
 atgccattgc gagagctctt gatcattgga atcattatct tgcggctaatt cactttatat 240
 tggattc 247

<210> 62
 <211> 442
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 62

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 aacgcgcgtg tcttatagcc tataggaggg tgagagagga ttcttccatt aatagagttg 120
 acttataaca acaattctca ctctacgatt catatggctc cttatgaagc tttgtatggt 180
 agaaggtgta ggacacccct atgttggata aagcccggag aaggccttac cttacgaccg 240
 gaagtggtag aacaaaccac cgagaaagtc aagttaatcc aggacaggat gatgactgct 300
 catagtaggc aaaaaagtta tcatgataag aggaggaaag atctgaaatt tgaggggtggt 360
 gatcatgtat tcttgagaat cactccgtgg actgggggtg gtcgagcatn gaaatcccga 420
 aagctacacc ttactttatc ga 442

<210> 63
 <211> 371
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 63

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 accctttgta gagcattatt cccaagtgct gtaagaccaa ctgtaaaaga aaaaaatta 120

aacacttgac aatggatgca tgcactacta tcaactatacc agctagcttc attcgtctct 180
 ttcaagcatc tatagcaatt ctttgcaata naatcttgaa actaacactt ggacagctag 240
 atctaaccgt tgttgctcga gtgtgaccaa attaatgggt atatttatta tgaataattg 300
 aatattanaa tactcttggc agtgcatacc tacanagctc acttgtggga caaaacatta 360
 cggtctttaa t 371

<210> 64
 <211> 442
 <212> DNA
 <213> Glycine max

<400> 64

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 accacattgg tctctgcgtg tcatcggaact tggggtctcc gaataacgag gtgcggataa 120
 ccgtaaagtg ctctgcatgc catcgaactc ttgggtcgtt ggatagcaag aaggtgacac 180
 taaatagtct cagtcggaag acgtcacag ctccaggaag agtgcagatt accacattgg 240
 tctctacgtg tcattggact tggggtgtcc gaatgatgag gtgctaataa ccgtaagggtg 300
 tctccgcatc ccaccggact cttgggcccgc tggatagcaa aattgtgaga caaaaattgt 360
 ctcgaccgga agatgctgac atctctgtca dgggtgcaga tgaccacatt ggtctccatg 420
 tttcatcaga cttgggatct cc 442

<210> 65
 <211> 376
 <212> DNA
 <213> Glycine max

<400> 65

aaaattacta catatcatcg tatgcttata cctacacgca tttttgcaac ctccaattgg 60
 ctctatagcg gaggtacaaa aactgcatca cgatcagctc actatacgaa aactataatt 120
 caactcaatc ctgcgaatg tgtatggatg atctacgcag gagcatgaaa cagacctcaa 180
 cacacttgca tgatacttag agaacagtgt gttgcgcaaa aagtctgcgc tatacttgcg 240
 ccttaaacac aaggcgctat atccactgaa cctgtactgg ttagaccacc acctacggct 300
 cgtagctcac acaactttag agcacctatg ttgagtccta tccaccata acgcacgctt 360

aagaatatct taagcc

376

<210> 66
<211> 370
<212> DNA
<213> Glycine max

<400> 66

tgcaggatga ttgggtttta ctttatgctg atgacgctac acaggatagc tttggtataa 60
gactgggctt acttgtacaa accatcgagg cgagatccta tgggttgcca tgagcatggt 120
gaaccattga tatcgaacac acatggcata agctgtagct gtcacatggg ctttatagat 180
agctagcgac ctcttttttt tttttttttt tgacacttgc ttggaaacgt atcgcttgaa 240
cattgttttag acttggcaca aactaggaa gctttgacct tectatttg aaagcctcct 300
tgacgattgt tgacagctgc acagtataag aattcatagc ctcaataagt gttttgctca 360
gcatacagga 370

<210> 67
<211> 378
<212> DNA
<213> Glycine max

<400> 67

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gggtaaattg ggtgttgagg ggtcaaactt tgaatcggtg gagttttcgc cttacaatca 120
ctttgagcaa gtctaaatta atgttatata ctggtttgag atgagaattt actccaaaat 180
taccatttc tcattttcac ttctcaaacc ttgaaaattc actcaattaa tgggttttgg 240
atacctagat ttggatttac cttgatctga agctggtttt tgcgttaaata acaatttata 300
catgatttac gacttgtagg atccaatttg agcaaaaatg gatgtgggca agaattggatt 360
cgaaatctgc cctattat 378.

<210> 68
<211> 296
<212> DNA
<213> Glycine max

<400> 68

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 catacatcat gtctcgccac ttggaggcct tttgagccaa acattaactt ttggccataa 180
 ccttggccta agatggaaat ttccaacctt accctccgaa gagagaacaa acgaatcttc 240
 ccaaacgaag cttctttttac cttgagttat aagtgtcgag ccagacaacc gattag 296

<210> 69
 <211> 365
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 69

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 gtaatgagga ttaatgtata agacaaatgg agtagagccc agcccaaata gttgaaatga 180
 ataaagtaca actaaggctc tcaaggntct tactcaatat aacccttaaa cactctntga 240
 gcctttctga tcctttcttt catagccttc gtaccctga ccacgttaca agcccaacaa 300
 agcccatgtg gatcaaggaa ggactaatta tgcttttgag tttggattct ggaatagaac 360
 ccaca 365

<210> 70
 <211> 321
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 70

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 tgtgatgtgc attgagcaga tcttcattct acaacagatc ttcttatact ttagctattg 180
 attacttctc atggcttaaa gttttactct tcatcagtga gcaatttgac ttcttcattg 240
 cataacataa ccagagacac tttcgagccc tcttttaagc attctcgcca atgatacttt 300
 ctgaatgact catgacaaca c 321

<210> 71
 <211> 311
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 71

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 actacttgaa cacaacatgt gcttatcgtg gatgagttct agggcattcc attgagcatt 180
 gtangaccct gaagcataag gtgcaaggtc taattgatgc gaggagaatc gcttgtgatt 240
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 aatgactcat t 311

<210> 72
 <211> 423
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 72

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 taaacatttt taacttgtgt aatcgattac acaatacttg taatcaagta ttgtgttattc 180
 gattaccagt gtttctaaat gttttaattt tcaaaattca aaatgaagag ttacatctgt 240
 tgatgtgtgg taatcgatta ccagtgactg atttcgaaaa atacatttcc aaaagtcaca 300
 attactcaag tgacttgttt ctgaagattc tttcaaaagt cacaactttt taagtgacta 360
 gttntaaaga aattgccaag agtcataaac tntgacttga gttatcaaga gattataagt 420
 atg 423

<210> 73
 <211> 397
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations

<400> 73

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taaatacacc cctgcccttt ttgtgtgatt cttttttcgt aaagttacgg aaacttacga 180
atttcgtaac gatacttatt ttctttccgt aatgttacag aaccttgcgg attacataat 240
catccccctt ttgacttacg gaatgttacg gaacctcact atttgtgcaa cgatgcttcc 300
ttttgatttc cgggtgtgtca cggaacctta cggattgcgc atcaatatat tcttttgatt 360
tccgcacgtc acgaaatttc acaaatngcc taatgat 397

<210> 74

<211> 465

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 74

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ttgagattnt aacagaaaca ctcggttaagc tgccaactaa attgtctatt ggtcaaccta 120
cacactcttc tgttttgcag gttatagggt gtaccatctg tgggtgaggct catgaaatgg 180
gccaatgtat tcccactaaa gaaaacactc aagaaattca ttatatggga aatcaacaac 240
gacaaaggta tactcaagga ggattttcag gcttcacgca gggccctat aatcaacaag 300
gacagtggag gacacaccct gncaaccagt tcaacaaaga ctagaatggg ctttcaaaca 360
gtccaatcca acaagggcct aacatattca agaggactac taagctggag gagaccttga 420
ctcagttntt gcaggttaaca atgtcaaate atanaagcac tgagt 465

<210> 75

<211> 418

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 75

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aaccgctctg cttctttcat caagcctgct gcttggtgct ccacaatatc ctatggggca 180
gcattggttg taataaaaag gtaaatttaa aaagaaataa atggtaaaga aaattatgtg 240
aaaagacaag gccactgaac caattcaaga cgactcaatt tttagacatg acacatgatg 300
gtaaaagtcc aaccttatat agcacaaagt tattacaagt tgcactgagg tttgataaat 360
ccctctcccc ctttcccgan aatgtataat caattacttg atttaaataca cttatctc 418

<210> 76
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 76

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tcgtttctcc gggagcgacg cgtccagctc agggacgacg aatatactga cttccaggag 180
gaaatagggc gccggcggtg ggcactactg gttactccca tggccaagtt tgaccagaa 240
atagtccttg agttttatgc caatgcttgg ccaacagagg agggcggtgc tgacatgaga 300
tcctngntaa ggggtcagtg gatcccgttt gatgccgacg ctatcggcca actcctagga 360
tatccgttgg tgttgaaga gggccaggaa tgtgagtat 399

<210> 77
<211> 397
<212> DNA
<213> Glycine max

<400> 77

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ttctagcaaa agttacaaat aaaataagtt taaggacctt tagtaaaatg aaaatatgcc 180
ccatatttgg actgagagtg acaacagtat ggactatttt tattaacgtt ttgacctcaa 240
aaatgagttt tctatgtttg aaaatgtatg gtagcgtata atatttgtga gaatccgact 300
aacagagcac caagagcact aaacataagg tatgagcgaa actgtgaaga actgagtcac 360
aaagagattc tattaccgta gatgacttaa ctttggga 397

<210> 78
 <211> 461
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 78

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 aacaatgtgt gctttcatca agtaaattct tttggcatca tcaaaacctg cacgattcac 180
 atttatgtca ctcaacctct aggttttgag atcaaaagga aggaattaat ggtgtacaag 240
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<210> 79
 <211> 347
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 79

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 tttattttcg naaatcaaac catgggcttg gaagggaac cgccagctgc agattgtagt 300
 attctaagta gtggcggttg cacgatcata tagcagtgtg ggacttt 347

<210> 80
 <211> 445
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 80

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tagtgcagta ggcttgcgct tagcactgtc tgcaactcaa aattgttctg caatttgcgc 240
ttagcatgag atgtcaggct tagcgctaaa tcaagctcta acttacaggg atagtccang 300
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cgcttagcgc atcttgtcgc taagcccaat tcatgaaagt tcaattccag ggaggaaatt 420
gagcttagcg cangacagcg cgctt 445

<210> 81

<211> 387

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 81

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cggttctcgc ttaacttaaa aataaaatag acttccaccg aacgtgtgaa ttgattcata 180
tctgacatca tattgagttt ataagcatct tcttgaaaag ttgaacaagc ataagacatc 240
atattgtaaa tcaaacaagt actaaaacta tgcaaccatc cgtgtttcat actttcagta 300
tcgtgtttaa attatgatgc atatcanatc atcatgaaat tcttccactt ttgaaagcac 360
caatgaaatg ttctgtctcat ggtcagt 387

<210> 82

<211> 294

<212> DNA

<213> Glycine max

<400> 82

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aaacctcgct tagtttgctt atgagcataa actagagtgt gatgctacat agatttaccc 120
tgcaccacgc cattagttag atgagtatgc gtaaccacat agaaccactg attcggaaca 180

tgatatacat atgacgatga gttattctga tcctagatat aattaggatc cccgttgtag 240
atcgggtggct catgcacttt cgacttagac accaactatg gatgagtcga ttac 294

<210> 83
<211> 315
<212> DNA
<213> Glycine max

<400> 83

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gattgattct ttccttccta tcattctcac ccttggttctt tcaaaccaca attccagaaa 240
atccacctct gcccaaaatt atcttgtgac cataactccc atgttacaca ctcagattaa 300
gtgattcttg agcct 315

<210> 84
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 84

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aggtagcaat ggcaaatcat aacagcact 449

<210> 85
<211> 361
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 85

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caactggggt accaagttga ccaacgcac cagtttgctt tcaagcttct tagtttcaga 180
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<210> 86

<211> 344

<212> DNA

<213> Glycine max

<400> 86

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aaagaattcg ccaaggacta accgcctgaa ctcttgatgt gcctctcttc tcctttatac 180
aaaagaacaa aggactaacc gcctgaattc ttttgtgtct cccttatgcc ttgacaaaga 240
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tcaaaggact agccgcttga caattgtttt cgatcccat tcac 344

<210> 87

<211> 402

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 87

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ttaataaata aataaataaa taaataagta aaaataaaat aaaataatat aattaggtca 180

taaatttcca ctatataaat caaatgttaa cctagagcag cttttacaaa acacttatgt 240
 ccttttctct tcttctgacg cacaagaatc ctaacagagc aactggagga ggagctctag 300
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<210> 88
 <211> 451
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 88

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 cacaaaggaa gggtagtcct tgtgtgttta gaactcgtac aagaaattta caagatagtg 420
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<210> 89
 <211> 563
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 89

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 catacaggcc attcgagtcg gcaccgtga ttctgtatat ctgcctgnat gcatgcaagc 180
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 atactctttc cctagtcgat cactcactta ataatacata ttctcgccct ttgattttga 300
 gtttatgctt cacttcgaat tagatcaatt acttacgcga gtccttgatt taatccctat 360

THE **WISDOM** **OF** **A** **POOR** **MAN**

agcttgtaga attatgggggt acccatcaca tgtggtacta ggtggcgggtc gggcgatggt	60
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ctccaactga gctcacgtac tcccacgtag cccatatcct cgttttctctc aacaccgggt	180
ccccatcaat cctctcaagc ttccacaaca tccaagcaaa acaacattca aacagcacia	240
gctatcacag ccaagcaaaa cagagccaag gcagaaaact ctgctcaaca catcaaccia	300

aatcacagct tttctcacgt agagaccaca gtaacaattc cttcgatcca attcgttaac 360
cgctggatcg actccaaaat tatactggaa gtctat 396

<210> 92
<211> 360
<212> DNA
<213> Glycine max

<400> 92

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tatgcaaggc tagacactcc atctatccaa ggagctactc caccactggc taaatatata 180
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<210> 93
<211> 407
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 93

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ttatgtgcgg gataatcctc acttgctcaa gattagagca aataaattga tataagatgt 180
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tgcacatttg tatgtgacac aatatgacaa atgccacaga aaatgaggaa tttcaagaac 360
gaatgagatg gccttgaaca acattcttga agntgaagtt ttgactg 407

<210> 94
<211> 287
<212> DNA
<213> Glycine max

<223> unsure at all n locations
 <400> 94

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 tgggtggaaga atctgaagag atcctatgat gccctctagg gggaaggaaa ccatacctct 180
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<210> 95
 <211> 380
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 95

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 agaacatgac caaaacaaag ttntgagggg ctttataggg cagcaatagt gagctcaagc 240
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 ggcttgcctt angtcgaaaa gaaatttgtc ccaacagtta aagtgagact gaanggaata 360
 tgtgggcat catcgatgag 380

<210> 96
 <211> 432
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 96

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 cttgagacat gaatttgata agagtttttc agaacaaaaa ggtcttatcc tcttataaag 180
 caaaatcggt ttatcctctt acaaattcct tggccaaatt acttgtgatt caataaggaa 240
 ttatttgagt gctcaaattg ttcaatctat ctctttcaag agagatttct tcttttcttc 300

1. The first step is to identify the problem or question that needs to be addressed. This involves understanding the context and the specific requirements of the task.

agcttgcaga ctataccttc gaccaaacac ggccgtgttt ctgtctcggc ccggatttaa	60
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<211>      458
<212>      DNA
<213>      Glycine max

<223>      unsure at all n locations
<400>      98
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41

<210> 99
 <211> 349
 <212> DNA
 <213> Glycine max

<400> 99

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 actccttaga agggctacca tttatcaaaa ttgatatcga tgctgagtgg aggcaagctg 180
 atatccaaga tctccatttt ggacaaaaac ccattctgca cagcatgtaa tccagaaaag 240
 accaagaaac tgaatcgtag gccttttcaa agtccacctt aagaatcatc acaggtttct 300
 tatttctcct tgcttctcct accacttcat taaggatcag aataccatg 349

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 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 100

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 cagtaatagc aactttacct gaaagtgtaa actgttaatg cagcacccaa tccgccaaaa 300
 caatgggacc atgatcctct ggaactctac catgttaata acttccaaaa tttcttccag 360
 ttcaccaga aacatcacct ctttctgact atttgttgct ggccaatatt tcaacaagcc 420
 gcttatcaca at 432

<210> 101
 <211> 406
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 101

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 gaggtccaa gaggattggg ctagagctgc tgaagaaggt cctaagggtt tcatgaactt 180
 taggatagat ttttgagccc atgggccaag gttgggtcca attatctttg tacgtattag 240
 attacgatgt cactatattt ggttcttgta attagggttc cataatgtag gtaggggtacc 300
 ctagaaatat aggatttttc agcccttgta ttttagggca cctagactag ttnttgtatt 360
 aagggtagtt ttgtaatttc acatgcacta agtgaatatt taatgt 406

<210> 102
 <211> 451
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 102

tntaaggccc tcaaagtact ggatggcaac acccactta aatgattatg aaagagggat 60
 aaatatgtga gattcttgga caaattaaaa gaaggaagt gacctgtaaa gttattggaa 120
 gacaagtcta gataaacaag ttggctgagt tctgagaatg aactaggaag tgttccatta 180
 aattggcagt aagctagatc aattgtagat aactgcttca tattggaaat tgcacctgga 240
 agctttcctg agaaatttgt atagctaaga ttcattgtat gaagagaacc atgttgtggg 300
 aagtttggca aagaaccccc aagatcttgg ttgtctgaga tgtcaaggac cttcaacggt 360
 gatatttga atatatcttt tggaaaagaa ccattcaagc cacaacttct taactctagt 420
 gtgactaaat tggagaaatt acaaaggatt c 451

<210> 103
 <211> 410
 <212> DNA
 <213> Glycine max
 <400> 103

agcttgctgc tattcctgta tatgtgtact gagatatttt ccttgagctt tgatgcaaaa 60
 aatgatttat ttgcatgtta aaacatagat ttaaccttaa atttcacca aatcatagtt 120
 ttctagcaaa agttacaaat aaaataagtt taaggacctt tagtaaaatg aaaatttgcc 180

[illegible]

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<223>      unsure at all n locations
<400>      104
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<210>	105
<211>	411.
<212>	DNA
<213>	Glycine max

agcttcttca	gaccatgata	ctctaggacc	tcaggaaagc	aaaatgttat	gaattcaact	60
tgggtgaatc	aactcaattg	ttagatagtt	gtgccggtat	aatgcttcac	agtgtctctt	120
tgcaaaccat	ggcataaatt	tgggagggga	tgagtgtttt	ggagattccc	cctttgtaga	180
tcacccaaca	actttctttc	tcttcttctt	cattttctcc	tctatgagct	ttgttttctt	240
ctctttttta	ggcttaaggt	taaagggagc	attgttgatt	gcaaccctct	taatatgttt	300
ccttctaggt	tgggtctaaca	tgggtgatgg	gaagaagaaa	gtgatggggt	gaaggaatta	360
cggaagaaga	aagggatcgg	aaaaaggggt	acttagcatt	cccaaaaact	t	411

<210> 106
 <211> 463
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 106

ttgatctacc accaccgccg ccaccatcat tntagttntc tattatttaa ttttactagt 60
 actttgtttt ctagccgtgt atttggctat attatgacat ttggataatt tagtatttct 120
 ttatttgcat ggtttgattg aacaattatg aattatgtta tatgactatg tggtttttat 180
 atatttgatc tattcatgtt tcttccttca tgattggctt atattcttca atgtatgtct 240
 tgtgaatgat taatagtata tgtttgcct atacttgta cgcactttgg ctttttggtg 300
 atgccaaagg gggagagaaa tagggattaa atcaagaact cacataagta attaacttaa 360
 tttcaagtga agcatanact caaaaaacac aggcggagaa tttaagtga tgttcgacta 420
 ggacaaaatg tgtgtatgaa tttcttgatt tcagggttat cat 463

<210> 107
 <211> 406
 <212> DNA
 <213> Glycine max

 <400> 107

agcttgcatt attgatggag aaaaggggaac aaccatgaat ggacaaagta atttagctgc 60
 tactaactct gtttttctact taagcaaate aatttctagt gatttttaac catcgaaaat 120
 gaatcttata aaattactgc ttcacatca tttagtgggtg ctatttgatg gcaataacta 180
 aaaacattaa cggaatataa gtttaaaaga ctaatcacia ttttgtttgg gaactaaaaa 240
 taaaaacctg aagaaatcca gacataataa acataatcta cccaactta tccaaaatac 300
 attacattct atgttctata atcccttttc ctttttttaa gattttttta aaaatattat 360
 attacaaaaa gtatatattg ataaagcatt cttttttttc aatttt 406

<210> 108
 <211> 388
 <212> DNA
 <213> Glycine max

<400> 108

tgcgatgat ggtcatgatg actacggggg atgacgactc ctgtgaggaa cacacagctc 60
tactgagtca gagaacatct caagttattc tggatcatgt ctagagtctg tgattcaagg 120
agaattcgcg tgtccagaag aaagcctaga ctcaagaatc tagagtctct caagaatcaa 180
gatcagcatt catgactcat gattctagaa tgatgaatag actctttccc gatcagtatt 240
aacgagtttt tgtcgaactt tgaatagcac atgagtgttt gacagaacct ttaccagcgt 300
agtttgactc tatggcgctc aattaccagc agcacaatga gctcgaaaaa agttttcaga 360
ctgaatttac aacgctccaa atattttc 388

<210> 109

<211> 318

<212> DNA

<213> Glycine max

<400> 109

agcttatatg caatgtggta ccatgtcagt gaataacctc gtcgggcgcc taggagtaca 60
tgacaagaca aaccacacaa taagtagtca agtcactctc actaggtaat atcatagga 120
gaccagtcag ggtcacagtg ttttgcgaga atgatccaac catatgggat caacataggc 180
ttaaaggagc actcaaaccg tgtaaccccc aaggcctaca ctccgaagag ttcgtcaggg 240
cctctccctc ctgattcagg tccaaccag aaaaatttta gcacacagac tctatctatg 300
aactgtacda aacacact 318

<210> 110

<211> 165

<212> DNA

<213> Glycine max

<400> 110

tcttatccaa cgctcatctt ggtggagaag ctcttcttc catggcttat tccctagtgg 60
atggtgcctc ctctcactg atctactttg ccttccgcta tatctccatg gcggaatac 120
gccattaaag gacctcatcg aagctcatgg aaccatccta catag 165

<210> 111

<211> 394

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 111

tataaagctg tttctggttg tatttaagtc ctaagctata gcttctttcc tcatgtaacc 60
ctgttcagac ttgttatata tatatatata tatatatata tatatatata tatatatata 120
tatatatata tatatatata tatatgggtt tgctaatgca ctcacaccat ttttttagta 180
tgagtacatt agcaatcaac accaaanatt ttaactaaaa aatcaatcac ctctcttatt 240
tttcaaaatt taatgtaggg atatcctaca tttgttccaa gctcgacaca tctgatctat 300
atgctcgagc ttgaggggga gtgttgaaat atgataagtt cccattggaa atgatctgtt 360
cctaattcta ctactgtttc aatgaaatta ctgc 394

<210> 112

<211> 330

<212> DNA

<213> Glycine max

<400> 112

tagctggtcc ttgtttgctt ctccacacac ctctctctct ctctctctca tgcactccat 60
atatcggatc ctagatacac aaacaaagaa agggctatac catatgaggg atatgaatga 120
aagaaagtct atagcgcagt gtcatttttt tggattagaa atatttcccc aagctgaacg 180
agaagcccgc catgccacgg ctgtagatat ccgttttaat atgaaaacat tcccgtgggg 240
ctttcataag aacaatgagc ccttcataat ctatgagaga aagcacaact catcgaagct 300
cgatcattac ctatactatc ttgtttatat 330

<210> 113

<211> 229

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 113

gctttgatgc cagctatgag gtatgtgcat gtttgaacag acatttatgc ttcgcctata 60
ccccctcct tggtctctag cgcaagttac tcctttgaca agtactggga gctggagtaa 120
aatgatgaca tgccccttag ctggactgac agcgactaca gtctggacta tttttagtca 180

cgacncgacc tcttaaagga cctttatatg attgtaaatg cagggtagc

229

<210> 114
<211> 407
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 114

atgatataag actttgaggc caaactatat ccaatatcan aatgaggaat cttttaacaa 60
tttaactaag tccttatttg tgattttctt gatcttattt ggtgacttgg aacaatgtgt 120
gctttcatca cgtaaattct tttggcctca tcagaacctg cacgattcac atttatgtca 180
ctcctcctct aggttgtgag atcacaagga gggatttaaa ggcgtacacg tcgaataaag 240
ccttgatggg tcttagacca gcacctacag cctggaatat aaggattgac tcagtttaca 300
ttgacaatga cttcaccaac tacacagtgg aatatgatgt gtatgagaaa aggataacta 360
cggaataact tttgatctgc ctatatgctt atgatttgct tgtgact 407

<210> 115
<211> 378
<212> DNA
<213> Glycine max
<400> 115

agcttatacc agcccaatcc cccaaatttt ttaatccgag ctgggaattc tctcccgact 60
caagtaaaag gaccacctgc aacagaaaga gcgccggccc aacgcacggc tccagccgct 120
ccccggccag ttaataatac agcccccgac gcgacctata aatatgcaca gcacccgccc 180
ccgaaagata acttctcccc tattcccatg gcatactccg agttatggcc ttcattattg 240
gagaatcatt tggtggtggc catacccggy aaggtcttcc agccacccta cccaagtgg 300
tacgaccggg gtgccaagtg tgtgtaccat agtggagctc ccggacacaa tattgactcc 360
tgcacccggt tcaagtat 378

<210> 116
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 116

tccattgttg agttgttget tcccttgtca cgtcttaatt cactccccac aagtaagtgc 60
aatttccctt ggttatttgg ctctccattg atgtgttttg gtgcttttagt tgctcatttt 120
tttgcaaaat tctgaagca attcgcattc gaatccatgc ttgttttggg gaattgaggg 180
tttgtgtgag aaggcattat gcctatgttg tattctgaag caatgggggca tgccacattg 240
tccccattct cttgcaattt gagtccgaac gtgcgcccc caagtgtctg gtgaagtgcc 300
ccaatgatat atgaatatga ttttgacaaa ttgggatggg gggactgttt tatatatgta 360
gagacagcat aagagattca aaatatgtgc ccgaatgcaa tttcaagctt atgaaccan 420
acctttttatc ttcaatgcaa gaagacatac tcatagc 457

<210> 117

<211> 305

<212> DNA

<213> Glycine max

<400> 117

agctttgata gtaagttaa ttgaataaat tatactcact atcacaaaaa tggctttcta 60
cgacgcacgt tttacgacgg ttgtacaaaa accgatgtca taagtaaagt agtgacattt 120
ttgtaaataa cttaaaaatt ttaaagatgg ttcttatcaa accagtcttt gaaaaggaat 180
taccacatca gttcttctac aaccgacgta gaatgcgaag cttaaaaatg cgaacgggct 240
ctctctcact ctctattata tctcttttat aatctctctc ctctctaaaat ctagaaaacc 300
ctaat 305

<210> 118

<211> 427

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 118

tctctgcctc ggntgaaaca ttggcgggta catgtgacct catgtggcca cccaaggatc 60
ttccacaagg gaagctcttg ctgcagaact tgcacatgtg tttcacttct tgatcttctt 120
ccattggatt gatttttcaa gatcacacac acgcacggat cagcaaagaa agcaaaatta 180
accacacttt cttgatcacc accaacacaa gagaaatcga tcacaaggga aaaacagcaa 240

caccccagat cagcatcaca tcttgaaagt ggttggagag aagaataata ccgagaagaa 300
gaagaagaga aaccccatgt ctgaaaattg caaggtggtg agtgcaagat ctaacgcaga 360
aaacaagagg aaaagaaaag ggacaagaga acgtgtagta gtagcacaaa ctattatata 420
tactata 427

<210> 119
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 119

agctntttgg agtagaaaca taggaccaac tcattntatt tcaaaaagaa agtcgtatct 60
agtcaaggtc tgagagacca tacaagtttc ctaacgattt ctaattatgt gggccattaa 120
gtctatcata tgctgacaat agccgagaag cccatgaatc tcttcggggg cggagtaagt 180
gtctgccatc gccttggcct tggctaacaa gcgnggaagt tcttgactcc cgttcaaggt 240
aagagcaaac cgatccatcc acatggttgc ctcttggtgt aaagagtcga tcacccttcc 300
tctagcctct ttttccgcgt atacttgagc atactcgtcc gcgattctat gctcgtgggc 360
cgtggctaga cctaactctt cttggtactt ggcg 394

<210> 120
<211> 335
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 120

agctntntat tttcagtaga tgaagatgaa ttcgtggcca cctcatggac tcctctaagg 60
acaataacat cattttcttgc tctgaattgt tgggagttgg aagccatctt ctcaatcaaa 120
ttcctagcct caacaggggt catatcacca agagctctac cactggcagc atcaatcata 180
ctcctctcca tgttactaag tcctcatag aaatattgaa gaaagagttg ctcagaaatc 240
tggtggtgag gacaacttgc acacaatttc ttgaatcttt cccagtactc atacaagctc 300
tctccactaa gttgcctgat gcctgaaatg tcttt 335

<210> 121
 <211> 439
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 121

 ntaacctcat cgtctctcac agtcttttaga attgggagcc aatccaatcc ttgtgttcgg 60
 actctcagcc acttatgata gccgtcgatg atcccattac tgcttcccct aagctctatg 120
 tcctttcttc atgccgcac ccatgccttg cgaactcctt ggagtaccct cacgttgtgg 180
 tcaccgaaac cccgtgcgat gaaaggcgtg atgctttcgt ctgatggcac tcctctcatg 240
 gggtagccaa gctgtcttat ggtgaggacg ggattataat taatacaacc cttgtttcca 300
 tcaagggaac atttggacat ccttcgcgatg aagatagaat cctgattctt ccttccttct 360
 agcgagggaa caaattaata gacgcccctc catgctagcc aagagttggt cccaattcgc 420
 ctttcctttt tcgatgcac 439

<210> 122
 <211> 397
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 122

 agctttaagc caattcatac gacaataact ttntactcgg atgtctgatt gagtcccgta 60
 atataacgaa acgctcgaaa ttgaatggtg aagctctaac cctattcaaa caacaataac 120
 gttttactcg gatgtctgaa tgagtctcgt aatatatcga cacgctcgaa attgaatggt 180
 gaagctctaa gcctattcaa acaacaataa cgttttactc ggatgtccga ttgagtgcag 240
 taatatatcg agacggtcga aattgaatgg tgaacctatg agccaattta aacgacaata 300
 actttttact cggatgtctg attgagtcgc gtatatatcg agacgctcaa aatgaatggt 360
 gacctctgag ccattcaaga caatactttt actcgat 397

<210> 123
 <211> 427
 <212> DNA
 <213> Glycine max

 <400> 123

gtgtgatgat tatttgtgaa gacatttgtt gaaagcttgt tgaaattgcc atgtttggat 240
gagttagaca taccattctt gtttaggggt ttttgtgatg atgcttgtga tgtttatatg 300
ctgaaattgc ccatggaaaa ctgctagaga tga 333

<210> 126
<211> 405
<212> DNA
<213> Glycine max

<400> 126

agcttcttag tttcagatga tccagatggg tttgtagcta cctcatgcac tcttctaag 60
actatggcat catttctggc gctaaactgc tgggagttgg aggccatctt ctcaattaaa 120
tttctggctt cagcaggagt catgtctcca agggctccac cactggcagc atctatcata 180
cttctctcca tattactgag tcttccataa aaatattgga gaagaagctg ttctgaaatc 240
tgatgggtggg ggcaactggc acatagtttc ttatatctct ccagttactc atacaggctc 300
tctccactga gttgtctaata acctgagata ttcttctga tggctgtggc cctggaagca 360
gggaaatctt tttctaagaa tactctctta aggtcatccc agctc 405

<210> 127
<211> 388
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 127

agcttgctct atatttacat tgatgcttat gggaagaggt tgtatgcat ttttgtttta 60
agagtagtgt cccactggta aaactaactt tccaaatggt tgccttcgca ggaaatggcc 120
ccaaggaagc ttgctcata gaggtctagg aaggacaagg cagccgaagg aactagttcc 180
gctccggagt atgacagtca ccgctttaag agcgccgtac accagcagcg cttcgaggcc 240
atcaagggat ggtcgtttct ccgggagcga cgcgccagc tcaagacgtt aaagaagcgc 300
tactaggagg caacctagta ctttttaaata ttctgcctgc tatttgatca ctctttatag 360
tangacgcac ctaggtgctc atgacctt 388

<210> 128
<211> 458

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 128

 tgcctaatta acctagaatt gagaganaat gattattaaa cacaaaatat gaaaataata 60
 agtattttatt acctatactt aacagaaaat acttataaca ttacaaaata accataaatt 120
 gggagagttt gatacaattt atacaagttt tatacacaaa agttagtcac tttcaccgac 180
 taacaactcc cccaaattta cagttttgct tgcctcaag caaaaagaga acaactcact 240
 agtgctcgag tgacaatgac atgcagtgcac tatgtacaaa ggtgtatgct acaaagtgac 300
 tgattgcatg ataagagaat ggagtaaaat gccctaatac cttgtctttc acaaggtatg 360
 cagttatcca aagagaagaa taaattgtac cctgaacaga tagatgaagt taggaataag 420
 acagatatca aggaagtag cttacaccat agtctcat 458

<210> 129
 <211> 347
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 129

 agctttacta ttatcttccg aacaatatat gaatgaaatg atgaacctta taattaaaca 60
 aagatactac tactactaag tttctattga tgcttgatc tgagtactaa aaagaaagcc 120
 tgttataatg attcaaaggc ataacaataa acaacttaac aataaaccat gaactacagc 180
 agctggnggt actttaataa atctctttgt attttaaaat agtctctaaa attntatgta 240
 aaaaagataa ctttacttat atttactaac taatgatata aaactaattt gctaacgatt 300
 taaactaaaa caaaaatgag ggaccttaac aaatctcttt gatttat 347

<210> 130
 <211> 398
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 130

 ntaacctcat cgtctctcac agtctttaga tttgggagcc aattcaatcc ttgtgttcgg 60

actctcagcc acttatgata gccgtcgatg atcccattac tgcttcccct aagctctatg 120
 tcctttcttc atgccgcate ccatgccttg cgaactcctt ggagtaccct cacgttggtg 180
 tcaccgaaac cccgtgcgat gaaaggcgtg atgctttcgt ctgatggcac tcctctcatg 240
 gggtagccaa gctgtcttat ggtgaggacg ggattataat taatacaacc ccttggtcca 300
 tcaagggaac atttggacat ccttcgcgat aagatagaat cctgattctt ccttgcttct 360
 agcgagggaa caaattaata gacgcccctc catgctag 398

<210> 131
 <211> 312
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 131

agcttgatg ggtgagttat aaattataat acctcagctt ggttcaaaag atcatgttta 60
 ctccgtcgca cggtgagcaa gttttgcaca gcttgcccct ctatggacat cctcaccoga 120
 ctatgtctta ctctggcctg aactcgaacc aatgcctgca tgcaccttaa tgccncgct 180
 gtctgcttcc tcacctgtct cccccgaac aagtgttga atcctcacca ctggcttcaa 240
 tgccctcaaa gcccttcttg cctgcaatca ccacacaagt gatgtcgaaa tataatgttg 300
 gtctaattgt aa 312

<210> 132
 <211> 414
 <212> DNA
 <213> Glycine max

<400> 132

tgttcatgtg agagctcaac actaagttgc tgatgtcttc acaaagccct taatgcttct 60
 aactctctct tcctttatcc acaagttggg actcattaac atttactctc caacttgagg 120
 gggattataa gttgtatgaa gaaatggagt tagttacttc agatagaggg tagatagact 180
 agttggtaat taggtagaat gaagttgat actaagtttg ttaagctgga tataaaatag 240
 tgtgtatgca accttatatt caataatcat caataatatt ctacagattt ccttggtgca 300
 caaagctctc tatcaataaa ttcccctttg ccaagtccac attgaagaat ttagagcaat 360
 tgtagaatgt cgaagaacat attatgtaca tacaagacac aacttataaa tctc 414

<210> 133
 <211> 397
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 133

agcttgctct aaatttacat tgatgtttgt atttatggga ggaggttgta tgccattttt 60
 gttttaagag tagtgtccca ctggtaaaac taactttcca aatgtttgcc ttcgcaggaa 120
 atggccccga ggaagcttgc ctcanagagg tccaggaagg acaagacagc cgaaggaact 180
 agttccgctc cggagtatga cagtcaccgc tttaggagcg ctgtacacca gcagcgcttc 240
 gaggccatca agggatggtc gtttctccgg gagcgacgcg tccagctcag ggacgacgag 300
 tatactgatt tccaggagga aataaggcgc cgacgggtggg catcactggt tactcccatg 360
 gccaaagttg atccagaaat agtccttgag ttttatg 397

<210> 134
 <211> 450
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 134

ctgatagcag atgatatgat ccttactagg agtggatcgc ttgatacagg tcatagagtt 60
 tggatgatgc tacttccaga gagggaagat aagtcaggat agacaccaca agaattgcct 120
 tgataagtct gagattgggt caacatgaga cccagagaga agctctctcc aaagtttata 180
 aaaggccaaa agtacttata ttgaaaatga aacccatata tatagcgtat ctgaatgaaa 240
 aaaatataaa tagaccaggg ccttcanata agttagggcc aaaattacga caataaaatt 300
 ataaataaca aatagaacat attttgcatt ggctttcaaa ttagtttggg ctttcaacaa 360
 caattaatat tcttagtagt gcctctggct ttggaccttc atccttctcc acttgagccg 420
 tggtaagtat gtctgttacc aatttgtgga 450

<210> 135
 <211> 394
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 135

agcttcttga tatactgcag cagcattata ttaatgcttc atagtatttt ttgacatata 60
tattaatatt gtcaagctaa aggcacaatt tttgttatgc taattgataa caatgataga 120
aatgatctta gatacaccaa gctgccttgg atatgatggt tgattcgatg atgattgggt 180
agtcagagta aatgttagtt tagagaggaa gaattaacta agttaattat attaaagatt 240
aataatttga aatgagacta ttaaaacata tagataataa attgtcataa tgaaaagttt 300
acgatctggt tggaatatga taaaatagga taagatatat cattatgata aacttttaggt 360
ttatttaata tttgatgcac actanataat atta 394

<210> 136
<211> 440
<212> DNA
<213> Glycine max

<400> 136
ctataaggac atgcatccag tcaacacacc gctgccaatt cccttatgaa taccctgaat 60
cataaataag ttcttcaaac aatagtttaa gttatgactg aagtttcata tacgcaaagg 120
caaataataa taatagtaat tgagctaagt tttttggaaa gtagttcaaa tttaaaataa 180
aaaatattaa atccatatta aattcgtaaa tttgaaacat tgtaattgta cttcaagtga 240
cgattttttt ttttttttgc agcagggtcaa atcaaggctg gtatctttat tttgagccaa 300
aatttcttaa gcagcgtttc acccgcaaat tatgcaacgg cgaacaaaaa ctttagaact 360
gaaaatcttg cattccaagg aaccatttag ccattggcct caagacaaga actacgtatt 420
gtggctgggt tgatccctac 440

<210> 137
<211> 340
<212> DNA
<213> Glycine max

<400> 137
agctttcagc tatgtatctc attagtaata ctttttcgtg ctgttgatg ggcaacaatt 60
gcctccttgg tgggtatagt aatcactgtg ctttgcaata ctccactcgc aaagttacag 120

cacaagtttc aaagcaaact tatggtgaca caagatgata gattgaaggc ttgttctgag 180
 gctcttgtga atatgaaggt gttgaagttg tatgctggg aaaccaattt tagaagttct 240
 atagagagat taaggaatga ggagctcaaa tggttgtctg cagtgaatt aagaaaggca 300
 tacaacacct ttctcttttg gcctctctg agttggctct 340

<210> 138
 <211> 453
 <212> DNA
 <213> Glycine max

<400> 138

ctgcttgatg agaaagaggc acagttacgc caatttataa ggtaagcacg tgttggttta 60
 ttcagtgggt tagcatcttt tggacatgct tgggtgtctg attaatttgg attgaagtat 120
 ggaagggggg agtactgggt catcgagcaa tttgctgaa gaaactgctc gtgtgtcgaa 180
 agattcttca gagctttttc caactggcat tccacaagta ggtcaaacag agattagcca 240
 agattcattt gcgggtggac tggggaatat tcgttcggag ttgattggct ccacatctgg 300
 caatgattct actacttttc tatcgaatga ccgtatgaga aatggcagag ctgacaatgc 360
 cactctaaaa ggtcatgaca gctccattag aggcagacag agatatactt cattgctgct 420
 caccctcctt gtcaagcgat gttaattaat tgc 453

<210> 139
 <211> 304
 <212> DNA
 <213> Glycine max

<400> 139

agcttgctct aaatttacat tgatgcttgt atttgtggga ggaggttata tgccattttt 60
 gttttaagag taatgtccca ctggtaaaac taactttcca aatgtttgcc ttgcaggaa 120
 tggccccgag gaagcttgcc tcaaagaggt ccaggaagga caaggcggcc gaaagaacta 180
 gttccgctcc ggagtaacgac agtcaccgct ttatgagcgc tgtacaccag cagcgcttcg 240
 aagccatcaa gggatggctg tttctccggg agcgacgcgt ccagctcagg gacgacgagt 300
 atac 304

<210> 140

<211> 376
 <212> DNA
 <213> Glycine max

<400> 140

ctgaatgctc tattcaatgg agttgacaag aataccttca gactgatcaa cacttgcaca 60
 gtggccaaag atgctgtgga gatcctgaaa accactcatg aaggaacctc caaagtgaag 120
 atgtccagat tgcaactatt ggccacaaaa ttcgaaaatc tgaagatgaa ggaggaagaa 180
 tgtattcatg acttccacat gaacattctt gaaattgccat atgcttgcac tgccttggga 240
 gagagaatga cagatgaaaa gctggtgaga aagatcctca catccttgcc taagagattt 300
 gacatgaaag tcaactgcaat agaggaggcc caagacattt gcaacatgag agtagatgaa 360
 ctcatggtt cccttc 376

<210> 141
 <211> 402
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 141

agcttgctgt agtataatag gaagcaccaa tataaatatc ttgatgccta ctattgatat 60
 atagcattga acatacatta aactagctag agagaataaa aattgctgat aatagtacac 120
 tccataggta gtcattgatgc gtaaaccact actgcaagaa aacactttat attatagtat 180
 tagctagcaa tttttgtggt tggcttgatc atgcttggcc tcacatgact gacaggtggc 240
 gaatccatct gcctatatag acccttcnc ataacttcct ttnttactac ttcttaagaa 300
 aatttcta ataggaaacagt agaagaacat taccatgaga cttccatggc tgagaatgaa 360
 cggtcagtta aagagacgta ttgtgtaaat gtgttaaaat ga 402

<210> 142
 <211> 391
 <212> DNA
 <213> Glycine max

<400> 142

tgtaataacc ttagatagaa atgagagatt atgttcttga gttgaatccc tcatgtgagg 60
 tgaaaatctt tagcatgtgt cgtaaaacca tagatatattt ttttatgtaa agtccaatag 120

tgactagcaa aggtgaaatc cagtgggtgca cctgggtctag tagaagattg aagtctagta 180
 aggaattgac aaggttgtga aaccaatgg ttgctggacc agttgcgaat tggttgtggt 240
 actgaaataa catctttaac ggtgaggatt ggacgcaccc caaggggtgtg gtgaaccatt 300
 atatagacct ttgtgcactt tcttctctgt ctctatatatt tgctcttgca caaatctaac 360
 actacttttg tataaaatac tacaatttgt t 391

<210> 143
 <211> 378
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 143

ntggacaacc atatataaga agtactttca gattagtcac agttgacagc cactccggaa 60
 gcatttcaag attgtcgcag tctgaaataa tcaagggtccg taaggagttg gcagtttctt 120
 gaagccattg aggtaaggcc accagctgtg gtaagcccca gaatgcaaca tattttaacc 180
 ttaacttgag gttatgctct tcatggcggt ccttccacag atctaagtcc agactaacac 240
 agtctttaac agacagagat tctaattcag gaaaatttat aacatcctct gaccttgact 300
 tcagactatg acaggcagca acattcaatg ctttaagagc aggggaacttc acccctgcaa 360
 agatagactc catattat 378

<210> 144
 <211> 369
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 144

agctttacac aaagaacata gtatagggtta aataaatata gctgaagatt taaatcacat 60
 agaaaacacc tttaatttct agtaaacata gctaattctc ccaactgctg ctcgctctgc 120
 aattctattg caatactcta catggaaaaa tgaggatatat tatttgtgat ataaaatgac 180
 aatcaaact aatgaagaga aacgaagaan attagatacc cgagttaaca actttcacia 240
 attaaacaat atctncttgt agacatatta catgagcata cctttgctac atctagnttt 300
 ccaagctgta ttgctagatc anatatgtag tcagggtcaa tagctacttc aagagcatct 360

tctatcata

369

<210> 145
<211> 397
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 145

tgtccacaaa nataggttnt tgaagtttgt catttcaatt tctcactaag taaaatggat 60
cattttcaag gtccaatgcc ttanaatgat cacctcttaa agtaaaaaag aatcacttga 120
taagaaagaa ctacgtaggt cttatttctt catcgcaatt gaggaatacg taggagcaaa 180
gggaaacacc cttgtcgact acaaaaagag aaaaatataa aaaggggtga aaggatataa 240
ggacttaaaa gggaacataa aaaatcaagg tcatgtttgc acattcgatt aaagcctgcc 300
gtcccttggg acggacgtgt ggngtgctaa taccttcccc gtgcgtaaac acaactcccg 360
aacctttcac ttannagttc gtagatcgcg tcttttc 397

<210> 146
<211> 416
<212> DNA
<213> Glycine max

<400> 146

agcttcaata cgagtagcca ccatttcact agatagtttc ctaaccagtc aaaccattcc 60
caaaatcatt ctgcagaata ataaatgcaa aatagagttg gctaaacaaa aaagatcctc 120
ataatcattt cccaattggg tcttggattc ccatagcact atacaaatca ataatttttt 180
aacagaaagt atactcaa atgaaacctag caattttctc aggttaaaaa aactacacct 240
cctagcaatg gagaggattg ctctacatc aaatgcttca gctggatcgg cctggattgc 300
acgtagctct tcattggttt cactagcaac cttatacata agccagaaat tatttctaaa 360
gcattgcttt ccatcacagg caacaacaaa attaaacatt atacaaaacc gagaat 416

<210> 147
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 147

ntganagagt tatcttttga caacttctaa ctctttttcc tgtaattntt aacacgtgtg 60
cccttacttt agtgtaagac ccattaaaaa ctatttcaca aaaattgcat ctaacttcaa 120
aatttccacc gcttccactt agacctttaa gctttgaaac acaattccat aaagggttgg 180
tgtcatcacc ttgttcttta acttgattag aagtactcat ctctacaatt aatcaatata 240
ataaaataat aaataagaac acaatggcca attttaaata aaaaaattat gaatgggttaa 300
ctgttatatt taaaaactat tatcaaaata ggatataaaa ttaaaaatca taaagtatta 360
aatgtattag taggtaagaa agtaaagaan aataatatta aactaaaaat tcttatgagc 420
ctacgaaaga agaanaaaaa attataaaaa ttggaaa 457

<210> 148
<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 148

agcttcatgc ttaactatgt atggaaaaac ttcattactg ttgttcaaga catacaagtg 60
agcttgtaac aaatcttcta cacttggagt gatcacctgc agtcctcttg aacccttacc 120
accactctg tcatcatgcc gacactcagg aagcccaaca gctttagcct tctctaagta 180
ttctgaacaa aattcaatgg cttcttctgc aatgtacctc tcaacaatag atgcttccgg 240
acgatataga ttctttgtat acccttttaa gatcttcatt tatcgctcaa cggggtacat 300
ccaccgtaga taaacaggac cacagcattt gatttctctg accagatgca caatcaagtg 360
aatcatgatg tcaaagaaag cangggaaaa tacatc 396

<210> 149
<211> 462
<212> DNA
<213> Glycine max

<400> 149

tcaccaccaa gacagtgtct tggataagaa gcttagagag gatgcttcaa tagaggaaga 60
gaatgagaga gaaagagggg agggcggtgg aattgatgaa ggagattagg gagagaagtt 120

66504304

gaactttgaa gtgtgtctca caagtttctc attcatcaaa gttatgagaa gtgttacaca 180
 tgtttctatt tatagcctag cacaatggaa gcttccttgg gaagctaggg gaagaaagct 240
 tccttgagaa gctagagggg gctactcaca cctctccaat agctaagctc accccatgtc 300
 aagatgcatg aaaatacaat gggaaacttc cttgagaggt aaggtagctt ccttaggaag 360
 caaggaagaa agcttccttt agaagataga gaggggctac tgatgcaatc ctaccccaca 420
 agggcattgg atagaagaat ccaagtagat tgggctagag at 462

<210> 150
 <211> 406
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 150

agcttttttt tacggttctt attaaatccc atcccccttc ataagctaaa atactctttc 60
 atatacagtg atatggtcag acacaaaaaa aagtggaaaa tattcaagtg taacacacat 120
 tgtatttgaa aagcaacaac tagcttttaa gtacaaaaca agcaaaacta aaaattacta 180
 cttgctagta gttgaaactc cctagtaatt agcattaaca ttggatttat tcaaattcac 240
 ctaaatacaa catactaatt aacacgaggg tagcgtatta tctcaaakat taaaagtggc 300
 ataaaacaaa ttagcaagac tatnttagtt caactgcttt ctcaaagtca actctaattgc 360
 attctcactg ctttcgttaa taaacttggg tcaatgctag atcact 406

<210> 151
 <211> 440
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 151

tatcttggac ctcaagtcca tgtagttgaa cctcagcttg ctgaactcac ttcccaaccc 60
 atagctgaag atgtccttgc aacaattagc acttccagta ttccatataa aatatctcct 120
 ttaaccacct ttgcttcaac ggggtgtgtc aaagaaagga ttcaagaaat gttgtgcctt 180
 caacaggcca acctagaaga aaccatggaa aagagtttcc tcttttaatg catggaaaca 240
 aatttgctta tcaagaacac cttggaagct tcaactcaagt ccttcaacga ctttgttttg 300

ttcaatatga atatgcta gcaacaacga ctgcctccaa ccgccaacc ccttctgcca 360
ccaccagcta cttcttccac actcacaact tctaaaaanaa ttgtgctgtg cctcgacctt 420
caccacctta gtcaccacca 440

<210> 152
<211> 395
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 152

agcttgtact cttgttgagg gaacccttac tcaagctatc ctaaaatgaa tacaacaaac 60
cataaaaatt tcatcacact catagcaaat cagaaataaa tggcgtgaat ggtttatcat 120
tacagttaca acaaatcaaa cacaagtggg gcacaaatca aacatgaatg gtgcgaatgg 180
tttttctttg caactactat ccacanagtg agatagaggt tagggattaa acaaatcana 240
gcacacaagt aataatgtta tttgtagtgt gagaaaataa caattgaaga atgctaaggc 300
gcgaanaaaa ttcaattaca cttctccann aaatgagata aagcttcaag attaagaaaa 360
tacaagtgha aaatttaaaa gtataatggg atctg 395

<210> 153
<211> 376
<212> DNA
<213> Glycine max

<400> 153

atagcttgct cattgactct cgattgctac acagaatgac caagatcttt acggtgatct 60
gcagaagagc atagaccaca gactcttgcg acatgtgtag atttcttatt catgggaaga 120
cgatgtacta cggtgaccaa cggatgaagg tctccttcag gctttttatt ttcacttgac 180
gaggaggaat gtgcggccac ctgatggact cctctaagaa caatagcata atttcttgca 240
ctgaattggt gagagttgga agccatcttc tcaatcaaat tcttagcttc agcaggggtc 300
atatcaccaa gggcttcacc actggcagca tcaatcatac tctctccat gttgctaagg 360
tctcataga aatatt 376

<210> 154
<211> 264

<212> DNA
 <213> Glycine max

 <400> 154

 agcttagaag aattaaaaat gaaaaaaaaa actataataa ctaaaattgg aaagacgtcc 60
 acttataagg actaaaatta gaaaaataaa cttatagaga ttaaaaatta aaaaaaatgc 120
 taacttacag ggacaaatac atattttaagc ttaaaaataa cattattcta aaattaaaat 180
 ttgggctcct agttagcatc aaaacagtcc atttattaca attaagatca agccagagat 240
 acttaaataa aataaataaa aaat 264

<210> 155
 <211> 314
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 155

 tgttacaaat gtggtaaacc ttatagaaac taaaggctgn tcgagtcgtg tggttgtgcc 60
 ggataaacat ggctttgcag cagccaacaa aggctcgcaa tgacgatgga gaccagtgtg 120
 atgctcgtg gagatgcgtc acgcgatgtg tgcgtgaacc tcacacgcca aggccctgca 180
 tggggttgcg tgcggcgggtg ttatgagttg gctgcgcaca tggagtcgtg gacgatgtgg 240
 tggcctttgg cgatgatgat gggcattgtt tacctgtgaa aataaaaagt ggcaaggctc 300
 accacggacg cctt 314

<210> 156
 <211> 403
 <212> DNA
 <213> Glycine max

 <400> 156

 agcttaacaa tccttttgat ctatttcaaa atatttctat ccctatcata taacttgcct 60
 cactcatatc cttcatttta aagttacaag agagaaactt tttcgttgag catttttcta 120
 aattggaaat tgtgatgttg agcatttttc catcttaaat ctctctagta ctttattgat 180
 atatgctttt tgagacaagc ttaacaatcc tatgatctat ttcagaatat ttctatccct 240
 atcacaaaac ttgcctcacc catatccttc atttcaaagt tattagagag aaacttctta 300

gtctcatgaa gaagatcaag atcattagtt gcaagcaata tatcatcaat atacaggatt 360
 agaaaaataa ccttactcct actgaccttc agatatatac att 403

<210> 157
 <211> 450
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 157

gtatcccaag ctggacttat atacagtatt ctttgggtcta ccatatacta tggctctaac 60
 ctatcgacga acttattgga agaatccctc cggactagca tatatgatcg tcttaccac 120
 caacttctct ctaagctcta ttagtcgtta ctctcttgat ttacgggatg ttgtgagcgc 180
 cttgttcttt gacttgagta gaagtactca tctctacaat taatcgatat aataaaatac 240
 tcaataagaa cacgatgggc aattttaaata aaaaaaatta tgaatgggta actgttatat 300
 ttaacaacta ttatcaacat aggatataag attatgagtc ataaagtatt agatgtaata 360
 gtgggtaaga aagtaaagaa tgataatatt aaactaaaaa ttcttatgag cctacgatag 420
 aagaangaaa attataaaaa tcggaaagta 450

<210> 158
 <211> 401
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 158

agcttccaag aatcaagatc aagattcaag actcaagatt caagaatcaa gagaagactt 60
 aatcaagata agtatgaaaa gggttttttaa aaaattgagt agcacatgga tttttctcac 120
 aacatgttta tcaaagagtt ttactctct ggtaatcgat tactagattg ttctaatacga 180
 ttaccagtag caaaatgttt ttgaaaaagt tttcaactga atttacaacg ttccaattga 240
 tttcaaaaag ctcttatatg ttttggtaat cgattaccac tgtctttgaa cgttgaaatt 300
 caaattcaaa tgtgaagagt cacatccttt cgcataaag ctttgtgtaa ttgattacac 360
 tgatttggtta atcgattacc agtgattggt tctgaataaa t 401

<210> 159

<211> 387
 <212> DNA
 <213> Glycine max

<400> 159

tcttatccaa ggctcatctt ggaggcgaag ctactttctt catggcttat tccctaattg 60
 aaggcgcccta ctctcagctc ttctactttg tcttcgctg catctacatg gtggaaaatc 120
 actattaaag gacctcattg aagctcacag atccaacctg catagagacc ccacaggcaa 180
 gcttccatca taaccactct atttgccta ccagggatat ccaacttgga cactgcactc 240
 gccaaagtaca tacacgacat acatcattac aatgacacta tcaacatcca cagcatctaa 300
 gtctgatgac actatgatca tctacctgat cccgtctcga tgcattctc aacatcaaca 360
 gtatctgac tcaatgacat aatcaac 387

<210> 160
 <211> 411
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 160

agctttccat gaatcaacaa aatgaataga taactcacia cactattggc atccctgcct 60
 ccacaaagca gaagaagtcc atcagagcgt gcacttgcag ttgcatacct agtcaaagat 120
 taaatggtaa aatgaaaatc gcaccatgac aagttgcacc aaatgatagc caactaatat 180
 atttaaggta agtctacact tcacagaatc gattgtttac atctttcta atgtaagatatt 240
 taaagtattc tcttgtgctt acaaaaaaca aaaagcaata atgttgtaac tttgttccac 300
 aaacagtaat attaattgcta atcaaaatgg ccaataaga tgagattcaa ttcanaccct 360
 actcaagctg cagtctaact caagttntcg tacagaanat canaagaaaa t 411

<210> 161
 <211> 442
 <212> DNA
 <213> Glycine max

<400> 161

tgtagactga atctatacca taaaaaagat ttgtgttcag tacaccgctt atcataattt 60
 tttttttctc tagccattat ttttagggag gtagttagc taactcacgg attttaattc 120

tttacttgca tgatgattca tttttccttt ctatacacat tgtttttttt gaatgatttt 180
 tacatatgta ttgaatcaaa catttcagaa ttatcattta tattaacgct atttatccgc 240
 atagttaa at tggccatcgt caaacttaaa tatacggcag atatatatta taactttttg 300
 tataatacat gatttttaac aaaatcttta tttacatttt cttatgataa gggattagaa 360
 cttttttttt gtggacaaga gatacaagtc tctcattcct ggaacatata taaagctgaa 420
 tgaatatgaa atgccctccc gc 442

<210> 162
 <211> 324
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 162

tcaagcttca agttagaagt tgaacacttt tatgttagaa gtgtcaaag aggaaatctg 60
 aagcgttggtg ttttctatgg ggaactacaa ggcgctagga gtagatgggt tccaatctat 120
 tctctataag agcaaataag atgttggtgg tgaaaccttt tgtactctta ttagaagtat 180
 nttttataat ccactaaca ttgatgattt taattatact cttattgctc ttaattccaa 240
 gcaagatgtg ctactaaca tgaagcactt taagcgcata atgttggtgca atgggttcta 300
 cagacaatga caagaatatg tctg 324

<210> 163
 <211> 379
 <212> DNA
 <213> Glycine max
 <400> 163

ctaaacgatg ggctagctta agctagccag gcaactttca tgttcttcat tagagaaact 60
 agctcagaag tgtgtcccta atgatctagc ttaagctagc ttggccactt gcaaattgtg 120
 tacactgttc tttcaatgat agctttaaat atctcttcaa agagatcctt actgtagttc 180
 ctacaaagag actgaacgac ataaaccacc tcacagctag tacactaggc tcttaaaata 240
 tttctctaaa gctgagttta ttcaaagatc aaccaattg tgctcaaaca atgttcagaa 300
 gcatgagaaa catatcatag ttgtcacaaa aatcgcaaaa aacaagtaaa agaggtaatt 360

ataattgata tctaactct

379

<210> 164
<211> 173
<212> DNA
<213> Glycine max

<400> 164

tgtttctata ctttatacaa gaatgaagct ctgataccac ttgttagaca agtggcctca 60

catatattaa gaaggggggg gggggtccca tcttagacac ccgcttcatt ctctcctcat 120

ccttatccaa aagtatttat tctattctac gcctcccata tattgatctt ccc 173

<210> 165
<211> 377
<212> DNA
<213> Glycine max

<400> 165

tgaaggatgt aagattctgt gattcttcaa tgtccaccac aatgtgatca aactttgatg 60

tcagtgttct gagcaccttc tcaatcacca gttgttcctt aatttgttct ccacagcact 120

tcattctaatt ggtgagtgtg agaattcttg tgaaatactc agctactgat tcagtctcct 180

ccattgcaag aagctcatac tgtcttctca atgtctgaag ctttaccttc tttatctttt 240

ctccaaagat aatccaatat ttatttagat caagatatct aaatatatat tttagtaaag 300

taaaagatag atacaattgc tttaatatat tatattgtta ttcttattct cgaaagatgt 360

tattatcata atatatt 377

<210> 166
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 166

tgttcctana tgatgggcta gcttaagcta gcctggaac ttttaagttc ttcattagaa 60

tagctagctt aaaagtctgc ccctaattgat ctagcttaag ctagcttggt aacttccaaa 120

ttctttacac ttttctttca atgatagctn taaatatctc ttcaaagaga tccttaatgt 180

aattcctaca aagagactaa acaacaaaaa ccacaaaaaa gcaataaaac taagttctta 240

aaatatttct ctaaagctga gtttattcaa ggatcaacca aattgtgctc aaacaaggtt 300
 cagaagcatg agaaacatat aatagttgca aaaaaaattg caaaaaacaa gtaaaagtgg 360
 taattataat tgattattta actctagtaa aaaaaaaaag cattgatcgt ctaaccttat 420
 tttatcaatg gttaaatact taattcaat 449

<210> 167
 <211> 331
 <212> DNA
 <213> Glycine max

<400> 167

agcttgggtct tgattttttt ctaagttctt taacaagatt agaacaatat acttgtcctt 60
 catttaactg tctttgggct tggcgggccac gatcaacaaa gtactttcga cacctactat 120
 atgttgattt gaccaacact gttatcggtg tggtgcgaca atccttcaaa accttattta 180
 tacattttga gaggttgggt gtcatgtggc catatcgacg tccttctcta tcataagcca 240
 tcgtccaatt ttcctttgaa atacgatcaa tccatgttgc tatggctgga ctgagttgaa 300
 cggaattttc taaattttga ttaaaaaaaaa t 331

<210> 168
 <211> 449
 <212> DNA
 <213> Glycine max

<400> 168

tgtgaagtgc cacgtagag aacagaacac tcttttcttc tctctgtttg aacaggctct 60
 tcttcacgag aaattattac tctctctact ttatcattag ctcttatttc agtgtttgac 120
 tttttgcgtt gttgcttcgt tgagtgtgca ctcggttctg tttggcgtct ggacaaaaaa 180
 catgggggaa gaagaagaag aagaagtgag cgtcaccgct ttgcatcatc caaacagcgg 240
 aaacgatgat cagagcctcg aattcgatat atatcctttg agcagttact attttggatc 300
 caaagatgct gttccctcca gagacctcac cttagatgat cgtgttctca ggatgaagta 360
 caagctcggg ttctttcttc tctctctgc ttatttctac tgcaatataa acactctctt 420
 tgttctcttc tttactatt ctttctttc 449

<210> 169
 <211> 394
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 169

agcttgtgct attcatcttt ntcattctct tctccctttg ccaacctgaa ttcttttgtg 60
 tctcccttct ccctttccaa gagaattcaa aggaccccg ctagaattc ttttgattct 120
 ttctttccc ttaaacaaaa gatttcaaag gactaactgc ctgagatatt tttgtttcc 180
 ccttacaag attcaaagga ctaaccgcct gagaattctt tgtcttaaca cattggaggg 240
 tacatccttt gtggtacaag tagaggatac gtctacttgg gttgttgaac taagaataag 300
 agaggggtaca tctcttgtgg atcagttcaa gtggagggtta catccacttg gttgttcaa 360
 gagaacaagg gaaggtacat cccttgtgga tctt 394

<210> 170
 <211> 463
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 170

tgtgctttaa gccacttaa caaaaaactg tttttataaa gtaaaaaagc cacactcttg 60
 acaagttcta catatgttgg caataaggtc caacatatag ctctctcatt tacagtaaac 120
 aaaactttaa aatttggttt aggcctcact cacccttggg ttggatcaat tgcaccatat 180
 aaatatgtgt tccccgtgga ctaacaaatt ttatttggtt tttatcgttg catttagacc 240
 tttgcatgat ggcgactttg atgtcataca cattacttgc gctctttttt ccttacacat 300
 ttgttgcatt tcttcaatct aatttggtga atctgggaat cagggttgtaa gttatgattt 360
 caaggaaaac agatttgccg atctacaccg atctgcgac agctntccag aatcaagggt 420
 cttctatgcc ggcactcctg ctacatcaaa tgcanaagca gct 463

<210> 171
 <211> 417
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 171

agcttgcttg tgggtgcttct atggaggctg gatctttgag cttcaatgag gtcctttaat 60
ggtgattttc tatcatggag atgcagcaaa agacaaagga gaaaagggtga gaagaggcgc 120
catccaatag gtaataagcc gtggaagaag gagcttcacc accaagatga gccttggata 180
agaagcttgg aaggatgctt caatggagga aaagaaagag ggagtgaag agagatgggg 240
gagcacgaaa ttgaaggaat aaaagaggga gagaagttga actttgagtt gtgtctcaca 300
agactctcat tcatcanagt tacaataagt gttacacatg tttctattta tagactangt 360
agcttcttg agaagctctc ttgagaaaac ttccttgaga agtttctatg agaaaac 417

<210> 172

<211> 461

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 172

tganatgagg aagtgtggaa ggggtgagact tctactttt attcggtggc cacagagtgg 60
tacctggaga tatgtcgcg nggtcaggag accttgggga cgtcagggtgg ggtgctattg 120
cccaaaacca agcttgacca atcccgaccc aaccgggca tagtcagtca gtgagaacct 180
gtgatgtacc taaacaggcg agctcctgga agtcaatcga taaaagaaca aagaccacaa 240
agcaaggagg cttgtgtggt ggctggccag ctgtgaatct tgagtgatat atgggatagg 300
gcctctggtg atcgattacc gaggggtgggt agtcgattac aaggcttana agtgaagaca 360
ggaagctaag atggcctctg gtaatcaatt accaagagag tgtaatcgat ttccaggctt 420
annaacgaga tcaggaagct aagagggctt ctggtaatcg a 461

<210> 173

<211> 260

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 173

tttcttctat gattcattct atgcacccat atatcgtgct gttggatgga caacaattgc 60
attctcggtg gggatactaa tcaactgcgt ttgcaatact ctactccat agtaacaaca 120

caagtttcaa agcaaactta tggtagacaca agatgatata gtgcgngcta gttgtgatat 180
 tcattagctt aggataaaat caattccgac cgttcggtcg tgccgtaacc acgttggaat 240
 tcaaagagag gtgaaaaatg 260

<210> 174
 <211> 237
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 174

tagtagaaga tanacactgc tagggacgaa cctggcaatg aagacatgct gtctactcta 60
 tatcggaacc ggtgataggt tcaaacctaa ctattttatt gatgataata tctagcagga 120
 cctatggcta ccatggaagg atgcagtgat tgtgacacta ttatgaatac tgatatggat 180
 tgtcacaatg cacgataggt tgtaatctat ttggacatta tcacgaagag gaattaa 237

<210> 175
 <211> 324
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 175

agcttgaaat tgaacaacgg aagctctcga gaaaatcgag tggtcataaa ttttcacaca 60
 gatgtccgat tcggggaaat aatatatcga gacgcacgaa attgaacaac ggaagctctc 120
 gagaaatttg aatggtcata acatttcact cgatggttcg atccggggac ataatttata 180
 gagacgctcg aaattgaaca accgaagctc tcgacanatt agaatggctc taacttttca 240
 cgcgaaatgtt cgattcgngg acataactca tctagacgct cgaaattgaa caacggaagc 300
 tctcgagaaa tttgaatggt cata 324

<210> 176
 <211> 381
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 176

ngccgccacg gagttntccg actatgctct tgtgtgttgg aacaagctac aaaaggagag 60

agcaagaaat gaagagccaa tgggtgatac atggacggag atgaaaaaga tcatgaggaa 120
 gcggtatggt ccggctagtt actcaaggga cttgaaattc aagctccaaa aactaaccca 180
 aggcaacaag ggggttgagg agtatttcaa ggaaatggat gtgctcatga ttcaagcaaa 240
 tattgaagaa gatgaggagg taactatggc tctgatttctt aatggtttga ctaatgatat 300
 ccgcgatatt gttgagctgc aggagtttgt tgaaatggat gatttgcttc acaaagcaat 360
 ccaagtggag caacaattaa a 381

<210> 177
 <211> 276
 <212> DNA
 <213> Glycine max

<400> 177

agcttattgg ttaaaaaggc gttatttgaa tcaacttagat aaaggtggaa gttataactt 60
 cctccattat taattaatct tctttcaccc actctctcca tatataaacc caccctcaat 120
 gctttcaaaa agtttaaaaa ataattataa gaaaaatggc aacatgtaaa cttccaccta 180
 cctaagttca gacaaaaaac acccacacac acaaagaagt taaggaacaa attggaacca 240
 taagcattct ctcttagtgg agaaggagaa aataat 276

<210> 178
 <211> 459
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 178

tannagttga aaataatata aaagtcctaa tacagggtcca gcatgattcc actcactagt 60
 acaaaaactca cctactaaaa tgtgtgctaa ccagtgtatc atacagtggc ctccattcaa 120
 tagttagtga aatcttttta ctatatttgt tgagtagtct aactaataaa tttccccatc 180
 taacctaata aacaaaaaat ggcaaatca ttttaagagca agcagagaca tatagagaac 240
 aagaaaactc agcatctcat ctaagacaaa atgcagcagc acctaagcat aatgcttatt 300
 acgggacaca tgagatatct caaggccctt ctgaataagg ttgtgaacat cctccanagc 360
 tacatcaactt ttttccttta tgaaattgca caacacacat aaaaagttag atttcaaccc 420

aacaaccaac tccatctgga tacaagccat gagtgaaaa

459

<210> 179
<211> 330
<212> DNA
<213> Glycine max

<400> 179

agcttgtaat cgattaaact gatatgagac atttgtctgc aagcttcaaa cacttggtgta 60
actggttact atcagtctgt aatcgattaa aacagaagag atgtaactat agaggaaatc 120
ttctaacttt agaacttttc ttctaactcc aacatgatga tgcattgatac acatatgaaa 180
tgatagagac aaagatgcaa cacacagtac aataatcaat acaaagtca ttcaagagag 240
ttgggcatgt agaagacaat aagatcaagc ttttctttaa gctgtaatgc taagtataca 300
tggtgcttcc cctatctcta acatgcaata 330

<210> 180
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 180

taatcttaca aagctctntg aaagaagggtt ttctatgtat tgattccttc aagtatccat 60
cttgagatac ttttcttgaa gaggaggaat atgcaattgg agtgtatgga gaaagggaaac 120
tagtgctcaa gcaatttccc atcagactca ttattgctcc taacccaaac agcctgatat 180
ttaagtatag gccaaactgca atgaagggtt atccccttgg tcccaccatt tttgtctttg 240
tgggataccc taacggaact aaggatgact atttctacaa cacatatgag gaaaaagtat 300
ctatcactaa gattggagtg ttccttgaga agcaatacat ctcccatgga gccaatggag 360
gaatgtagat cttggagaaa tccaagtggc acataacatt gatccaccta caatggaaca 420
agaattgatg ccacaaaagc attgtagatc catcttt 457

<210> 181
<211> 368
<212> DNA
<213> Glycine max

<400> 181

agctgtggaa agtgttgttg tcaccttctc gctaagccac tctgatggct tatcgagcgt 60
 ctgctatatg caacattcct gggctaagcg caaggaagaa tccataagaa gatgagctgt 120
 acaagtgcgc taagtgcacg cgcttcatct tactaagcgc accacttgag ttcattctgct 180
 aagcgagaaa agcgggctaa gccaaaaatc actaacgtgc gctaagcggc ccataagtgc 240
 gctaagcaca cgagcacaaa caaggccgcc tagttaagcc tgaaatcaga tcttgtgaaa 300
 ggagtatgga ctaagattca gagctctgca tgcctagggt ttctagagag agaaagtgca 360
 agttctag 368

<210> 182
 <211> 377
 <212> DNA
 <213> Glycine max

<400> 182
 gctattacgg acctataata ctacagctgtt tatccgtttt tgtgcaagac atattttaaac 60
 cgatcaattg tcattttaagg cgttggacca ttaacgatct cttgggtttt taaaagtagt 120
 ggtaaaggta gacgtttatt gtatgtttcc gaaggtgcat attaaccaat aaaagcagag 180
 agaacctttt aaggcattgg accttaaaac ggtttttagt gacttttgcg gacaaaagct 240
 tcatttgaga gttgatttta gccttaagtt cactttgggt attagtcaat tcattcaagg 300
 aaacttgcaa agaaaaatgc ccgactgagg ttttttcttt ttgagattgt attcaaagat 360
 attgcgatta ttttatt 377

<210> 183
 <211> 364
 <212> DNA
 <213> Glycine max

<400> 183
 agcttagaat ggccgaaagg gacgagtcaa ggggtgtaag catggcatta aaggataaat 60
 tgaaggcttg tcataggtca aagagaagtt tgaccgaaca atcgagtgga acggaagaga 120
 atatgttgac gatcattgat cagtataagg agaaggtaaa cctagctgct agtcttaggc 180
 agagactaga ggatgatcat gcgaaggat tgactctaca aatggaaagg gaagcaagag 240
 agaggggtgat agaataatta cgctgggaag ctgtgaaatg gatggataga ttcgctctca 300

ctctgaatgg gagtcaagaa gcttcagggt attagccaga accaggcaat gacggaagta 360
tact 364

<210> 184
<211> 439
<212> DNA
<213> Glycine max

<400> 184

gtctccacta agttgcctaa tgcttgaat gtcttttctg atggcagagg tcctagatgc 60
agggaagaat ttctccaaga acaccctctt aaggatcatcc cagttgaaaa tggacctggg 120
agcaaggtag tatagccaat cttttgccac tccctccaga gaatgaggaa aagccttttag 180
aaagatatga tcttcttgaa catcacgggg cttcatgggt taacaaacaa tatggaaactc 240
cttaagatgc ttataaggat cttcacctgc aagaccatga aacttgtgca gcaaagtgtat 300
tagtccagcc ttgagaacat aaggaacacc ttcacagga tattgaatgc acaagctttc 360
ataagtgaaa tcaggtgcag ccattctcct aagagtcctc tcacgaagtg gaggtagagc 420
catgttctca gtatgaaaa 439

<210> 185
<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 185

agcttggaga ggatgcttca atggaggaaa agaaagaggg agagaaagag ggagggggga 60
gcacgaaatt gaaggaagaa aaaggagag aagttgaact ttgagttgtg tctcacaaga 120
ctctcattca tcaaagttac aacaagtgtt acacatgctt ctatttatag actacgtagc 180
ttccttgaga agctntcttg agaaaactct cttgagaagc ttctttgaga aaacttcctt 240
gagaagctag agcttagcta cacacaacc ttcataact aagctcacct tcttgagaag 300
cttccttaag aagattccta aagaactaaa gcttaactac acatacctct ctaatagcta 360
agctcacctc ctgagatgag aactagagct tactcc 396

<210> 186

<211> 139
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 186

tcataagcat ctacccattg cgtcaanaga tacatgcaaa gtgctacaca tgctgctctt 60
 tatacactga ctacctacct tgatgggctt tctatgagga cactaccatg gaaaacccta 120
 ttgtaatatc taccttgat 139

<210> 187
 <211> 396
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 187

agcttcaaag aagactattg aagtgtggtt caatcaattt acaaactttt ggaggcagta 60
 gacaaagatg acagctggga aaggacaact agcagagtca ttctgaggta gtcattgttga 120
 ggaacctttc tgatgagcat tttaaagtaa aatcagaatg gtgatgttag ctgagaaagg 180
 atccatactt gaagtagatc cttttgatga agatcaacta ttatgtcttt ctgaagtagc 240
 agaactgcat caattntaat acatcttcac aacacanagc atcctgaagt agattgcttc 300
 attaaatcac agtgaaggca tagnttcttg gtgtttagtg gtctatcana agttagaatg 360
 tgtaaagtct tcaatcttcc attctgatga tccaca 396

<210> 188
 <211> 395
 <212> DNA
 <213> Glycine max

<400> 188

tgacactatc caagactcta tacaatactg aagctctggt ctctacagat cttcacacag 60
 cacaagtact cttaactctc tggagcttgt acctttctct ctctagaaac cctagacatg 120
 cacagatatg aattctaate catactgccc ttgtaaaatc tgaatatatg ctcatatatg 180
 cggccttggt cctgctcgtg cgctgtacgc acttatggac cggttaacgc acattagaga 240
 atttacgatt acagcgtgcc tttctcgcat atcgaatgaa ctgaatacgc gcacttaacg 300

agatgaagtg gtgcggtcac agaacgctta ccaatcaact tttccagag tcttgctcgc 360
acttaaccca tgaatgttgc gcttatcgga cactt 395

<210> 189
<211> 499
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 189

cgattgaccc tttgagcccg agatcccggtg agtcacctgc agcatgagag ctatgcagca 60
attgaaatgg tcacaactgt tctctctgtt ttacgagagg ggcacataag atattgagat 120
gctcgaaatt catctatgga tactcttgag caatacaaat ggtcagtgac ttttactgg 180
gaggtgagat acacgctcat atgatatcgc gatgctatac attgaacaac agaagatctc 240
gacagattca gaaggtcata tcctntcact cagaggtctt agtcaggccc ctagcatatc 300
gagacacgaa tattgagaga acgaatgggc tcgacaaatt catatggaga gagattttca 360
cttgtatgtc tcatatatgc gcataaggaa taaaaacgct cgggtctgtt gatggagagg 420
tctctagaaa acaaaggggc gttgtcttgc acgcatggca attcagcaca gagtgatgtg 480
acctcgata tactcttcn 499

<210> 190
<211> 428
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 190

gaggaagcaa ccctgctcgc ctgggagagc tgagctcgcc tgggagagct gggcggaac 60
cacctcccct attttgctat aaatagggga ggaaatgaag aaggaagggg tccggcccct 120
ttggcacttc tctctcttcc gaatttgctt ggaaaaattg tttccgtgaa gaaaatctaa 180
gccgaggcgc ttccgaaacg tttccgtaac gttttgcgtg aggaatctag cagaggtttc 240
aaccgttctt cgacgatctt cattcgataa gcacgatct tcgaccttct gcgggtaagt 300
accgccaacc aagctgttca atgagagtta tggacctcgt tggcttcaca ttcgcactcg 360
tgctatatcc tcttatagcg ctgacactag aacangaagc tgatttggtg tagcaagtaa 420

ataagact

428

<210> 191
<211> 238
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 191

aacattcttt tcagctaang attatatact tatgttcttt tcaaatgcan aatcactaga 60
tggatgaaca tgatgacatt gacatcttgg cttctagcac aacaggagaa atgttaacgc 120
ttttctccgt agggcatana aatcaatcta ttgcatggga tcacttttga taaatgccat 180
acccaatcc tttatcccaa gctaaatgtc gacattgtgg tgctntgatc aaatatag 238

<210> 192
<211> 210
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 192

tcatatctcc canaacccca tacncacgan nattaagaga gaaagaagtc caccagacc 60
tggattntcg aagtcccact cgtagccacg cacttcacga ccccgaaat gccctccttt 120
cgcgatttgg agcagaaatg agcaccaaag gttggagctn ntgtcgggtt tcaatggaga 180
atggaggaga aggaaaaagc aacgtgagga 210

<210> 193
<211> 220
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 193

gagtttcgaa acaagagtat aaagtgtgca gagtcaaatc tggatatttg cttggatgan 60
natgggctag aatatatcta anatcagata cggnnngtgg actgcatctg gngtcggaat 120
gaggganana aatatgecta actcaactnn taaaaanaat caactaaggc taccggagta 180
ctatacgaac tntggtcata tgctntgatg atgagagatc 220

<210> 194
 <211> 209
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 194

attattcaga tgctttatga atgacaatca aaggtgactt ggaaacaaga aattaaagag 60
 acttttcatt tgccaaacag ntntatcctc tcaaaaagaa taagaagtnt tctgaactga 120
 aatggtntat cctctcanaa agattctttg gtcaaccact tgcattattca ataanggaat 180
 ttgattgac ttcattgtac aatctatct 209

<210> 195
 <211> 194
 <212> DNA
 <213> Glycine max

 <400> 195

actactcata attaattgcta ctagaagagg atgactgata actattatta ctataatattc 60
 caacaagcta ctcaccattt aggaagttga tctgtattac ctgaacaatg gcaactatga 120
 caatgatattg ttatgttcaa acattataacc ttgagttagg ctatgctgta gaaatgagac 180
 taattaattgt catt 194

<210> 196
 <211> 366
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 196

accatggtaa tcagaataga aagtctactt cttatgtaag agttgcagag attcctttct 60
 cacttctcac ttattacagc tagaacctat gtctagctga tggttctatc ggctgtgatg 120
 caaagattca ctatcaccct gtatctaata taaaagtggg cagattcaga attgttactt 180
 tagaccacgt aaagaataat atgcatattt gacaatatag accaagagaa actcagatag 240
 cacaatccag tggtagagaa gcaacactta atttagtggt cannagtcgt atatagacat 300
 gagaatagat accttgccan aattatccat aagacattta tttccattcc ctaccagtgt 360
 attcat 366

<210> 197
 <211> 276
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 197

atgccctctc tctcttctcn tttctctctc tcttctggct ctcttatatt cgnnttatagn 60
 tctaggctct tcttagacac tnnnttcatt ntgcaattcc actnttagta ataaaaattc 120
 gctcttcaat ctataatttc gttctctatt gattaatgca aggctaagtc tccagcgtct 180
 gtttctcttg aggatcaagc acagttctct ctgaggtctt attattactg ggtaaattct 240
 gntcagtttt tctcttcact acatactctg aatttg 276

<210> 198
 <211> 234
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 198

caataatcaa taatctatct ttcaatcttc tctcaacatc attcaatatt tntcaactct 60
 ttctacacna atttctgatt catttctctt catctttcta aaagtttttg ttcaaacactc 120
 tctcttatga gaaaagttct ttgttcanaa acttggtgta ttcacccctt tcattctctt 180
 ctccctttgc caaaagaatg aagggaactaa ccgcctgaat tctttgtgct tctc 234

<210> 199
 <211> 499
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 199

atgaccctta gancttcaga acnctctaga gtccacctga ngtatgagag cttcagacag 60
 ttagcaatcc tgtgggtgcat tgttttatct gaactaccat tatcaagtaa tattggaacc 120
 accatgccat taattgtacc actaaacttc atagttccta ggccagagga acctttaagg 180
 gcattataag atagatgatg gtctaaattc atatgtgcc tggcttcctc aataatagaa 240

ggtggctctg gttgcacttc agcattttca tctcatcca ttgcaatagt aggcattgcc 300
 gattatgaca cctatgataa gaagtgaact tctcatgaca tgtataacat agccctttct 360
 cccttcgaat ctgcattttca gctgggtgaca ttctttctaac atttcctggg atcggaaaac 420
 aggagtttga tctaactcag tctaggcacc tgagtcacaca ttcccttgaa tggatatgct 480
 aagagagaga cttgtgact 499

<210> 200
 <211> 299
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 200

tgtcgctaga gctgacccat caactgtcct atctctntca gactggcgac tcttangetc 60
 ttgatcttga cttgatagaa cctctntnta agtgaagggt gcctgactcg atcccatggt 120
 tactaaagtg gaataaaaac cagtgcgaat caagactgtg acatctatca caggtgaaat 180
 ggatgaatgc ataaagaaat gcatatggca cagatgccat ttacggatac ganagccccga 240
 gagaatatct atttcttana tacaacattc nggcagcata gtgcccgatg catgcattt 299

<210> 201
 <211> 429
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 201

attattatct gaactaccat tatcaagtaa tatttgaacc accatgccat taattgtacc 60
 actaaacttc atagttccta ggccagagga acctttaagg gcattataag atagatgatg 120
 gtctaaattc aaatttccca tggcttcctc aattctagaa ggtgggttctg gttgcacttc 180
 agcattttca tctcatcca ttgcaatagt aggtattgcc gattaggaca cctatgagaa 240
 gaagtgaact tctcatcaca tgtataacat agccctttct cccttcgaat ctgcattttca 300
 gctgggtgaca ttctttctaac atttcctggg tntggaaaat tgganngtgt aattcatttg 360
 gggcttgga gtaacggcgg catgcttggt gacttttctt actgacacta ccgctgactt 420
 ctacttcct 429

<210> 202
 <211> 433
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 202

atgactattg aataatctat tcatgtttcc tttgatgaat ctaatgctat ttctccaaga 60
 aaggatattt tagatgatgt tgcagaatct ttagaacaaa tgcattattca tggacaagat 120
 tctaaagggg aagggaaagg aagcaatgaa gatcctccag aagaagccaa atcaaagat 180
 gaaggtggca tccatggagt tcaactgacta tgctcttatt tgggtgggact aacaacaaga 240
 agatttggag aacctttggt gaatacttag gaggacatga aaagattaat gagaagaaga 300
 tttgttcctt ctcattataa taaagacctt cataacaagc ttcattaggct catacaagga 360
 anaaaaagtg tagatggata ttataaagag atggagaatt ccttgagtag agccagtctt 420
 aatgaagatc aat 433

<210> 203
 <211> 240
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 203

acactcgngg atatgnggac tntctctatt caactntnnt gttcgtccta tcagtagaat 60
 tcgtttgtag ttccgtacac aattntcttt gcttcttcag ttacagtga gttagatatg 120
 ttctaattat taatttcttt tctttttttt tcgaatattc catntaccat aaatagttgc 180
 agaagcttct gtcacaccta taaaaacact ntatatttga cttgacgaat ntatatgtat 240

<210> 204
 <211> 437
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 204

gtgtccccc tcgaaatgan aaacctttat tcaaaccttt canagntagt gagaaggcta 60
 aacgaaaaat aggggaactta gaaaagctaa atccttaact gaaggcgtag gtgacaatca 120

tagtgaatta ctaaacaaga atggtagttt acttaagggtc attccagata cctcccaagc 180
ctcggaaaat acttctaana tggtacaag aagtaccttc aaatttaata atggtattaa 240
tgaagatagt gaccaaactc agatacacac ttggatagga cactatcaga aagatataat 300
ccaataattc aaaactgaaa cacctcaaat atatatcacg tcaactgccct gcctctatag 360
agaagaggaa acatttaaag ttagtgaata cattacgatg acataatgac aacgagatac 420
atatgatacc tcacata 437

<210> 205
<211> 505
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 205

ntgaccctga accctggatc tctaagtcac tctgaagatg cacgcntang aganaccata 60
aaaactaagg tagtttctaa actaaaatca attgaggaag cttcgccaag tatccccatt 120
gaaaaacctt tattcaaacc tttaaagtt agtgagaagg ctaaacgaat aattagggaa 180
cttagaaaaa ctaaactcctt aattgaaggc gtaggtgata accatagtga attacttaac 240
aagattggta gtttacttaa agtcattcca gatactcccc aagccttgga aaatacttgc 300
aaaatggtaa caagaagtac ctacaaatta attaagtta taattgaaga tagtgaccaa 360
agctcagata acacaactga gataggatca gtgtcagaaa tgaatataaa ttcaattaat 420
tccgagcact gggaaacacc ctccaaatta tattatcaac gtccaactgg ccttgacctt 480
ctattataag aaagaggaga aaacg 505

<210> 206
<211> 424
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 206

tgctgactcg agagacaagg acaaccact ctaaagcatt tacgccatca gactcactgc 60
ttgagggggtt gtacatcatc cagaaagcga atccaataa taccgactca agagacaagg 120
acaactcatg cttaagcatt tatgccatta cgcttaatgc ttgagggggtt atacaccgta 180

caagatgaat attctattaa taatgactca ggagacgagg aagactcacc cttaagcatt 240
 ttaggcacaa ggataaatgc ttgaagagtt gtacaccgct cgagatgagt attctggaga 300
 tattgacctct agtgaggaga tgacttatcc ctanacattt atgcgacaag gctgaatgct 360
 tgaggngtta tacgccattt atgatgaata tcccananat accgacctaa gcaaaagaga 420
 cacg 424

<210> 207
 <211> 511
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 207

ntgaccctga aaccctcgna gcacctgaga taccctacag acgatgagag ngatgcaagc 60
 ttccncaaca tccaagtaac tctacattct aacaacacaa accatcacag ccaagaatac 120
 agggcaaagg cagataactc tgcccaaaac accaaccaaa atcacagctt ttctcactta 180
 aagacccag taacaattcc ttcgttccaa ttcgttaacc gttggatcga ctccaaaaga 240
 ttactggaag tctctagtac ataagcctac attatgaccg ttgggatcta ctagcaaaca 300
 tccagaactc attctgaact actctgtcca cagccaatta cacacaagca ttgttctgca 360
 cttgtgcaaa attctgtctgc acaatttcac agcataaatc tgcacaaagt gcagatttctg 420
 aataccacac ttctctcat ccaatcttgc ccaaatcaaa tactacaagt cccnatcatg 480
 tatcaatcat gtctaaacca gagccaagct g 511

<210> 208
 <211> 257
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 208

actattccaa tacgctngga gcagcttaat agatctcaag ctgtgcaaatt attcagagct 60
 aatagattga gctgagtcta aataatgaag gaacagttat tcaagttagc ttcatatcc 120
 tccgggattc ggaaggacac attttgcatt cgatataatg attcacatcg tgatgtagtc 180
 ttgaattatc gtacatctat attacgccg ttctctattt ttgtacatga atatagacag 240

caacatgcaa tgaacag

257

<210> 209
<211> 488
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 209

accattgaa tcttgagacc atcgatananc cagagatcct ctagagacga cgtcgacgca 60
tgcaagcttg ggctgggcct acctccatcc ctagagttag ccgtatgagg cggaagctcc 120
acgtacggtt ntgaagccga gcctttctag caatggngcc tagggaccga tatgatgatt 180
ggtttaggta gggcgccgg cctactacgg gcacctgtag ggattagtgc ttgagaccgc 240
gateccacaaa agcatgggac tcacccttta cttgagaatg aagaggggaa tgacacgacg 300
tttcaagagc tatgcgaggg gtgaaccaa ctgcagaggg atcttcctga ccaggcgtga 360
tagagatgcc ctttattacc caactcatat tatcattcaa tcttgctttg tgccactcag 420
tcttgcgggg atccccctcc tttctctttt ctctcaaccg gcgtcccttt cctccacaaa 480
cgtgctcc 488

<210> 210
<211> 180
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 210

tgcattcttc ttcaactcct gaagccctat catagcagac tcaagaacat nctgcattcc 60
anagatcact ttcattngaa tgcactatgt attggtagtc ttcccatcca ggaatggtag 120
atgcgttgaa ggtccgttcc cattcatctc tgaaacactg aagcttcagg gatccacac 180

<210> 211
<211> 510
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 211

ggtgccctta ggcttgagac cctgtagan cccgtgatac tttgcagacg acgcccangc 60
 atgcaagcta tgatgatatg gtcttcacct tcgaaatgat caaagtgggt ctgagaagag 120
 gcaaactctga tcatcttgct ttgatacatg cgaacaaaca aagttggggc aaatacagag 180
 ggtgatgatg aatgagaacc ccgagctgtg actgacattc ctatacagcc gagtttccca 240
 ccaacccaac gatgtcatta ctcagtcgat agccaacctt ctcttacct accgcccagt 300
 tatccacaga ggccatccct ataatatcca cagagtttgt cgttcgcact ctcaatgacg 360
 accatcatct ttagcacana cctagagcac caaccaagat atgaatttag cagcgagaaa 420
 gcctgtagaa ttaatcccat tccagtgtcc tatgctgact tgctcccata tetaattgat 480
 aactcaatgg tagccataac cccaaccacg 510

<210> 212
 <211> 425
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 212

acatttgctt gttgctcgtg tgctgagtgt gtgatgagat cgtgtagagg ggtttgatgg 60
 aacttatatt tatattagtt gagtgtggct gctagtcctt gtttgtaggg gatattgtag 120
 tctttgcagt taattctcgg cttgtagata ttaatcaata gcttacatat aatgtttagag 180
 ataaacattt gcttatagat aaaaaggtag aagataatca aacctttag ataatgtgtg 240
 ggcttataaa taattatttt aactgccaat agataagata ttcaaataca tttgaatatt 300
 agtaggttag agataacctg tttgttgggg agtccggctg gtnacgttca tccctcctct 360
 cttttggcct gccttcattg tngttgcctc ctctcatga ttcttttctt ctacccctcc 420
 ttccg 425

<210> 213
 <211> 190
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 213

tgaagacccg cacanacatt ngaaagaatn tcacattgtc tgctccacca tganaccccc 60

agatgtccaa gaggatcaca ttttctgaa ggcttttctt cattcattag agggagtggc 120
aaaggactgg atgtattacc ttgctccaag gtccatcacg agctgngatg accttaagag 180
agtattctta 190

<210> 214
<211> 224
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 214

atatannntt tttagtatta tatagaattg aatgttngaa ttttaattgat cctgaggctn 60
tggaatgtga ttagcaatct aatgcataa agtatntat ttaggcgggtg ggtgatcaag 120
tcaatctctg caaactgttc anatttcaat gagattntgg gaaaagaatc tgagctgggtg 180
gaatanatag ctgaggctta atttcaatac aggaacatta tact 224

<210> 215
<211> 156
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 215

tcttggatgg ctactcgng tataattaga tagcagtaga tcctaaagac caagaaaaaa 60
atggccttac atgcccttcc ggtgtcttta cttacagaag ggatgccatt gngttatgta 120
atggccttgc caccttccaa agatgtatgc tagcta 156

<210> 216
<211> 195
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 216

aataccagaa gaatccaggt tctgaattcc atatcttgat tagaaggaaa ggtagcataa 60
tcaagtaaag caaaaaggga ttgaatttga ctattttctt tattgctcnn nnnnctngta 120
ttgtttgatt gtcttaaact aattctcttg tttcaagcat ggcaattaca ttcacaactt 180
actcataatt tatat 195

<210> 217
 <211> 181
 <212> DNA
 <213> Glycine max

<400> 217

ttccaccatg gagatgcagc ggaagacaaa ggagaagagg taagaggcgg cgccatccac 60
 tagggaataa gccttggaag aaggagcttc accaccaaga tgagccttgg ataagaagct 120
 tggagatgat gcttcaatgg aggaaaagaa agagggagag aaagagagag gggggagcac 180
 g 181

<210> 218
 <211> 274
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 218

gcagctgagc ganggtccaa tcattaccca ttatttacct tcntgnctta catttcttgc 60
 accaacaact tcgtcacttt catanttgga aggggtgcta catggaagtt gagggcagga 120
 gaggaanaaa gatggatata attatgtagc gagtggactg aagggatngg tgtggattct 180
 tccatcgtgg tcttcacata nttttggttt tgcatacaat acattgttgg attacgaaca 240
 cctaaatcgg acgaccttgn tagctcttca cata 274

<210> 219
 <211> 249
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 219

accatcacga ttatcgtctt ccttgtcatt attgggggta ccacctgagc cgccagatcc 60
 ctacaccttt tgagcgtggt ctttgaaaga tccgtcctcc tttgtgcaca tgctcatgag 120
 gtgcatccta tccgaaacca tatcaagatt gtactgatac tgactaacac aggcaaccat 180
 tatgtccttg caagaatgga ctctgtaatg ctccaagtta gtgtaccang taacagctac 240
 ctgagtaag 249

<210> 220
 <211> 246
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 220

 gtatctngtt ntccatcgat gtgccatcat tntcttctat gttctanacc ctgtctcgca 60
 ccatttaatt attgattggt ctttaattgtc aattaattag gcagttctat tatttgngcc 120
 cattcagcca atgtgatgct ttttaatctaa tttcaggaat taatgaagaa ttgngcttga 180
 atctagcatt gngcttgaat ctagaattgn gctcggactt gaagagggca aactatatta 240
 ttctat 246

<210> 221
 <211> 510
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 221

 gtactgcgcc tcgtacttca cancnctgta gtacccggca tccttagagt cgacctgcgg 60
 catgcaagct tgctctanat nttcattgat gtttgtattt atgggaggag gttatatgtc 120
 cattttgctt taagagtagc gtccactgg taaaattaac tttccaaatg tttgccttcg 180
 caggaatggc cccgaggaag cttgcctcaa agaggccag gaaggacaac ggcggcgaaa 240
 gaactatttc cgctccggag tacgacagtc accgctttag gagcgctgta caccagcagc 300
 gcttcgaagc catcaaggga tggtcgtttc tccgggagcg acgcgtccag ctccagggatg 360
 acgagtatac tgattttcag gaggaatatag ggcgtcggcg atgggcacca ctgggttactt 420
 ctatggncaa gtttgatcca gaaatagttc ctgagnttta ttccaatgct tggcaacaga 480
 ggatggcgtg cgtgacatga gatcttggcg 510

<210> 222
 <211> 207
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations

<400> 222

tctcatgcac ttagtgcca nataatat ctatagttag aaatttatta tggtatggta 60
aaaatagggtt tgttcaatct ctgtaaacca tanttgcat gntctttcaa ttaactgaaa 120
taatgagtct gtgagacatg acttaagttt aattctcaca gaatacactc ttngaaaatg 180
atgattagct ntaagtgtga ctaagtc 207

<210> 223

<211> 431

<212> DNA

<213> Glycine max

<400> 223

tctggttgat gagttatcga cagcgatgac tgccaacagt cctgagaatg tggtcaggac 60
tgattacgtg agatgcaatc tagaggatc tgatctatta gaagacttg ttcatcattg 120
cttgagacta tgacctcgac tatcctgaga acattctcgg aagacttcgg aagggatata 180
ttgaaatgag gtttcataac tccttcatgc cattgataag gatctcgacg agcttctttc 240
tctgtcttgt acgtcacgaa agatcctacg ctctggcgat cggtcaccag aaacagggca 300
tcggatccca ctggcgatca ttgctgtact attcattttc gtgcgctcaa gacgcttggt 360
tctgcattac cagaggtggt atcagaacgc gtcattcgca tatctctgcc actgtcctct 420
tggtagagaa t 431

<210> 224

<211> 481

<212> DNA

<213> Glycine max

<400> 224

ttggcctgga gctggtcttg agcaccgccg ctgcagcttg ccagcatct ggatcacatt 60
ttacgacgca atttggttgt gccatttctc acgactaagt tctctctgc tccttactca 120
caccacgaac aatatagttt agcgattaca ttgtaagagt attaggaact tgaaaactaa 180
gcacaggcta cactttaact aatggccttg ttgctctgc cattcttgga acctcagcct 240
actaattgaa acgcttcgag catgttggtt ttgactgcc atatgggggc gtttttctta 300
taaaggaaaa aataattggt tatataatcc gggggccttg atattgacag actgctccac 360

actaactttt ggtttatttg aaccctcctt tatagggtct aacacttgac acacgcccc 420
aataaggcct tgagaccaat atatgctttc ctctatccta tgggtgcctta tctcctatac 480
g 481

<210> 225
<211> 277
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 225

acctagacta taaatagaag catgtgtaag actaggtgtg actgtgatga atgaaagtct 60
tatgagatac acttcanagt tgcacttctt tgectctgtt attccttcaa tttcgtgctc 120
cccccttctc tctntctttt cctccattaa agcatcctct tcaagcttct tatccaaggc 180
aattcttggg ggtgaagctc cttcttctt ggcttatcc ctagtgatg gtgcctcccc 240
tctctcttc tcttttgct tctatcgaa cctacc 277

<210> 226
<211> 279
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 226

acatcccact gagatgcact atgtaagtac ctttcanaaa acagacttgt tgagaataag 60
ggagctagca tctntagtaa gtgatccagn tgattntcaa gctcaccatg ggaagttgct 120
cagaattctt agagtagatg ttgaggaagg atgcctagag accctgggtc agttctatga 180
cccgtctac cattgcttca catttccga ttaccagctn gtctcacac tngaagagta 240
ctcctaccta gttggcttac ctgtgccaga caagatacc 279

<210> 227
<211> 220
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 227

togtcttatt canaaccnca accaattatg anattcncta tctcccactt cacacctcgg 60

aacgcaccgt tcttatagag agaggcgctn tcacatcntt cttaggctgg gagaggaaat 120
 gttcccatnt tttatgatac tccgngaac agatatccag tggagatgac ggngtgngc 180
 ctgtagctca gaggattaga gcacgtggct acgaaccacg 220

<210> 228
 <211> 379
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 228

gcagcttaga tgatagaggt ggagactcag gtctttctct tgcttgactt ctttataatg 60
 tggaggactt ggctgagac tccgctagat ggggctatct gcgctacacg cttttctcgt 120
 tacgccagct gttccgctac cgcattggan gccttcaata tttctagttg attcttttga 180
 cttaatcctt tacgattgca acaggaagga aattttaatt ttacaatagc atgatactgt 240
 aatattttta gatatttata ttttagataa atactattta gagtatacac caagtgagtc 300
 acagtgtgaa tgactgaatc agtgtacacg acaacagcaa gtgaaatcgg tgacaacagt 360
 ctcacaaata tcggcacat 379

<210> 229
 <211> 169
 <212> DNA
 <213> Glycine max
 <400> 229

atacctgtaa tgctgcatag ccacagaaaa tctcttcttc tgattaacac caccttcgtg 60
 atcataagct tctggcattc tatactggca gttcaggaga acatatgagc cgttggaaag 120
 aaacatatag atttgtcaat ttacttatat atgcccataat ccctttgta 169

<210> 230
 <211> 497
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 230

nttaaccggt gagctttgag acccttcgag naccgtcatc cttggagaag agtctaggca 60

tgcgagcttg tgtaattaga tacaaatggt aatgtttatt tatataactg tgaaagcgca 120
 taagtgtggt gaaacaaaaa cctgaagtac catctcatta gtcagagtat atacaagcat 180
 gtacgggtga tagaaaaaaa aaagatgttg gaagggttga tgacttctta gtgcatagag 240
 tgttactcac ggtgatgatt tcttctcttg ccaccaaagtg gcatatgtcg gcgatggaat 300
 atatatgtta gaaggggagcc tggcaatgat ggcattatat tattgctttc acatagatag 360
 atatgtttta ataccttctg tgcacttccg tttatttatg aaatagtggg aggggtgttat 420
 ttcttgaatc cttgatTTTT tttAACgacc gaggtgacat tctatttttg tctatagatt 480
 ataacacttt atttaat 497

<210> 231
 <211> 503
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 231

tgggggttttc cggtttggcc tatagtcttg gacntcaagt gctctgaggt atgcagatta 60
 ccatcagacc acttncaggg tgctgtaact actttatatg gacttgatgg ggcctatgca 120
 agttgaaagc cttggaggaa agaggtatgc ctatgccgct gaggatgatt tctgcagatt 180
 tacctgggtc aactttatca gagagaaatc agacaccttt gaagtattca aagagctgag 240
 tctaagactt caaagagaaa aagactgtgt catcaagaga attaagagtg accatggcag 300
 agagatagaa aacggcatgt gtactgtatt atgtcatctg accgcatcgc tcatgaggtc 360
 tctgcactca tcacaccaca actatatggc gtatttgana tgctaacctc gactttgcca 420
 gaagttgcct ggacttcttt ctcttttaca acttcactca tttctggttt tttccttcca 480
 ccgtctgctt tttgatgtat tgc 503

<210> 232
 <211> 259
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 232

gtaatattat agccgatgct ctntctcggc gtcatgcatt actttctatg cttgaaacan 60

naatgattgg tcttgaatgt ttgaanaaca tgtatgaaaa tgatgaaact nttggagaaa 120
 ttttttaaaa ttatgatatt ttttcagaan atggtttctt tagacatgaa ggctttcttt 180
 tcaaagaaaa cannatgtgt gtgcctaaat tntctactag aaatttgctt gttttgtgaa 240
 gcacatgaag gaggttaat 259

<210> 233
 <211> 300
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 233

aatctcaata agtcaacaat gacattcgaa gaaatcgaaa tacgttatgg aggggatgta 60
 ctatagccaa tgtgtggaag gcaatagaaa gaatnggaaa ccttaatcta atgctaagtg 120
 tagagagtga gagagagaat gggaacttag taactcatga nagactntga agtctgaaca 180
 agttagtga tgtgttagca ctacatattg aagctttaaa tataaaanat atgtaaacad 240
 ataaacaaca atagtaattc taaagacatg tcacatcant gggcttatgg gttgggtcat 300

<210> 234
 <211> 271
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 234

aagaatgtgt atgtgtttct tgatttcagg gttgtcatca tcaaanaggg gaagattgta 60
 gaagcaagct tcatgatgat gaatcaagtt gattcaagta gttctgatga taacatagat 120
 gatgacaaaa agccaaaaga atgatntcaa gattgagtca acaaatataa gattaaattc 180
 aagaatcaag agtcaagatt caagaataat caagatcaag aatcaagact catagattca 240
 aanatcaaga gaagacttan tcaagataag t 271

<210> 235
 <211> 212
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 235

gagagagaga ggngaggnga gcataatatt gaaggaggac aagagagaga gaagttgaac 60

tttgatatgt gtctcacaag actctcattc atcaaagtta caacaagtgt tacacatgct 120

tctatntata gcctaggtag cttccttgag aagcttcttt cataagcttc cttgagaagt 180

tagagcttag ctacacacac ccctctaata ac 212

<210> 236

<211> 266

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 236

cacatcaaca tgctngaacc ttcacttttt gtgagnaata agatttatnt ataagaatgg 60

ngaacaagga tcaaactcta gaccacacag ttatagaggc tccaattaca tgtcatgaac 120

ttgntactca aaagagctgg tgagttaaga catatggatg aatttatgtc tagcattcat 180

gttatgggtg tatatttgaa tattgaatat actatatgct ntgagagggt ntaaacttaa 240

cttgctgtag aggaactgaa atatgg 266

<210> 237

<211> 338

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 237

acatctaact cagactaccg attcatgcca ataatatatc gagacgcttg anattgaaca 60

acggaagctc tcgagagatt caaatgggtca taactnttca catggatgtc caattcaagt 120

gcataatatt ctgagatgct cttaaatttaa catggaagca caagggaaat tanaacggcc 180

ataacctata acaaggatgt ccgattcagg ccaataatat attgagacgc tcgatattga 240

acacttatgc tctcaagaga ttcanatngt catacatnt cactcggatg tccgattcag 300

acgcataata taccaacatg ctgatatta aacatcac 338

<210> 238

<211> 269

<212> DNA

<213> Glycine max
 <223> unsure at all n locations
 <400> 238
 tgacccacgc ggggtgttgaa gagacggcat gggcatctcc ctcttcctt cttgcccctg 60
 atgccccgat tcttntggca ttcagcggtta tggaagaaac gtaatcaaac tttcctctnt 120
 tcaatccaac ctcgattctt tccccggcaa acaccagatc cgcaaagctg gacggcatgt 180
 aacctactag cttctcatag tagaactctg gcagagtgtc taccatcatg gcgatcatct 240
 ctctctcaac catgggagga gctacttat 269
 <210> 239
 <211> 246
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 239
 ctacactgac tcccagcttg tctaagggct agtggcanac atatatcaag ccaaggaggc 60
 agtattgctc aagtactatc atgctgcana aacccttatt gatgactnta atcgcttcaa 120
 gatgtaccat atacggggggg agaacatcac cagagcagac ttgctctcca agttggctag 180
 cactaagaga gctggacatc ttaagaccat tatgcaagag aactccaag cacctaccat 240
 agacac 246
 <210> 240
 <211> 271
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 240
 acaacctcaa gatttatcat atacttatca agagaatcaa tccttaagag agactntcca 60
 nattcattgg aagcactgag aatcttgaca aactgttaat atacagtaga tgtccttcta 120
 acagatctgg acatggatat gaaggagata cttatgttca tgataaggaa actaccaaact 180
 gttatttctg tggaaaggat caatgttngg tcaaagatcg cattggagac cttctacca 240
 ttgtcatgac ttgagtgcct tcaatcacta t 271

<210> 241
<211> 425
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 241

gatcgccgct atttcctggc cgacgccgac tgcattntt ttcgatcaat atcggtgaat 60
aatactttnt ttgccgaggt gggctaagt tttcctggcc gaataaatgg gaacatgcc 120
gtttcgggcy aaacgaaaca tcggttgagc tcgcacgaaa aaacctagcc cacctacatt 180
gtaagttttt tatgcaacac cgaaacaaga aaacttcccc tgccgtaaga aaaaacatta 240
tcggccagcg agcatttttt tttaaaaaaa attgcgcaat gtcggctgaa aaatatcagt 300
cggngccatt tcacgaccga tgcggctat tttggtttct attcaatccc tgaatgaaat 360
ttgcatgatg tcgattaaga aatgttngat ccgcgtgatc cggatgatgct tttttttaga 420
actcn 425

<210> 242
<211> 253
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 242

tcttctcaat gtccgttaag tcttatggac ttcttgccca taaaggataa gcctttttctt 60
cctgcctctc attnttttgc tctgcacgg ntgcacctcc aatatnttta atatccnctt 120
attaccctc tctgcttctt ctgttgggct tcttctccct tctcaacact atggcagaga 180
tcctatgcaa ctgaaccana ccanaatctt ccatcaagaa caaaccanat tngaaaaacc 240
acaatggaca aac 253

<210> 243
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 243

tgctttcttct agaagcaatc gccttctgga ggaattttct ggaaggccca agtgggccta 60

gttgctatTT gcaccccat ttttactaaa tacaccactt gctctttttc ggagatttcc 120
 ctccgcacca aaccccaact ctccccttgt ctcgctctct atcacgtcag cctcaaacc 180
 tttcttggtg tctctctcct cctcatccc aattactacc cctaccaac taaaacatt 240
 tacctcacgt cgcgtttact tcacatacct ccccgctcct gcgcattcct ctctaccct 300
 tcgccaccta gtccccattt tcctctntc tcctctcac ttctgcttc cacttcccc 360
 cccctcttc ctctctcct ctctctctc ccc 393

<210> 244
 <211> 416
 <212> DNA
 <213> Glycine max

<400> 244

tattttcctg aatcggacat ccgagtgaag agttatgacc attggaattt ctcgagagct 60
 tccgctgttc aaatttgaga gtctcgatat attatgtccc caaatcggac atccgagaga 120
 aaagttatga ccaattgaat ttctcgagag cctacgtcgt tcaatatcga gcgtctccaa 180
 tattatgcac ctgaatcgga catccgagtg acaagttata accaatcgta tttctcgcga 240
 gctttaggcg cgcattatac tacacctcca tacatatatt ccaactgactc gtcactctac 300
 actcaaacct attatcatct ccatactttt ttgataaagg ttctgatccc atactcactc 360
 ctatctctcc ttcatctcgt ccactcgttc ttattttctt tccactcgat ttatcc 416

<210> 245
 <211> 222
 <212> DNA
 <213> Glycine max

<400> 245

catgcttcta tcttgagatt atgacacatg ctaattcggg ggagatgatg attcgcgaga 60
 tgtatgcagt catctcagat attgtggttg gtactctctg aacaggtcat taagcgaact 120
 tagcatagtt agctctcttt tgctagagga caagcaaac tatacatact ggggagtttg 180
 atcattgacg tacataagtg gattatgcaa ttaagacata ta 222

<210> 246
 <211> 429
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 246

gggatgtgag ctcttgtaga cttttctttg caggaatgta agctttatcc aagaganatg 60

tatggtcatt gggtgtcatg ttttcaatta gctccatagc ttcattaggt gtttttaact 120

ttatcttacc tccaacggaa gcgtctagaa gttgcttcga gtgaggctcg aagccatcaa 180

taaaaatatt tagttgtatt ggctcactaa atccatgcat tggagtctga cagagtaaac 240

cgtggaagca atctagagct tcgctgagtg attcgtctag ggactgatga aatgaaaaga 300

tatctacctt tccttgtgct tgctttgatt ctggaaagta tttcttcaag aatttatcca 360

caacctcttc ccaagttctc aagttatttc ctttaaacga gtgaaaccat ctcttatctc 420

actggcagg 429

<210> 247

<211> 215

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 247

tctcactacc ctgtttctgc cataacttaa gcctatagat aattctttac cccatgtngt 60

ggctcctaaag aatagtataa tcatcacata ctaggtatcg tttggaatct tttaaagag 120

aaagaggata ctatctcgat aatcaatata taatcataat tcatgagtat caatctaadc 180

aatataatca taattcatgc taatcaatat atact 215

<210> 248

<211> 510

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 248

atgacactga gactctttgn actnccctag aatcctctag agtcgatgct gaggcacgca 60

agcgttggca aatggagagg aatattttgt ttctatatgg agacgacgaa gagagcaagg 120

tgttggaggt ttctgacaaa attgaaagga tttgatttga tcgcaagcaa taggtgtggt 180

agtaagggtg ggggacgaag gattggccgt gtgataggag ggcaacgaac aaatttccaa 240

ggcaacgaaa aatgatttac attttttatt ntgttcgcta aatgtttcat aagtcttact 300
 aaagtgaatg caagcaaaag aaacaaatat ataaaaagga taattaccat tttagtcac 360
 cttgaaattg caaggtgttg tcccattact ccatgacaag aacaatattc atattaattc 420
 acgtgtctct caatacgtcc ctacccttaa aactctttca tctcctcacc catcctgtca 480
 ctctctctct tttccatcta ttatgtgtcg 510

<210> 249
 <211> 239
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 249

atggtgtctt ggagcgtgct tgtgattntn tctttcgcca tgctgcacaa ctntcaggtg 60
 ttcccttgag aatggtggag agaagcagaa ggcagttccc tcttagaana gcccgatg 120
 ctgcagaaga catgtctctt gggctactca nagccaaggg tgatggattc atgacattga 180
 tcgaaaatgg taaatggatg tgtgacgagg cacctcacag tggaaatgaa tatgtaaac 239

<210> 250
 <211> 255
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 250

cgatggngca caacaagctn tccacatcca caatgcgcgc ataaaccac catccnctgg 60
 tgccacctc caactgagct cacgtactcc cacgtagccc atctcctcgt ttctctcaac 120
 accgggtccc catcaatcct ctcaagcttc cacaacatcc aagcanaaca acgttcaaac 180
 agcacaagct atcacagcca agcaaaacag agcagaggca gataactctg ctcaacacat 240
 caaccaaatt cacag 255

<210> 251
 <211> 230
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations

<400> 251
aatttctggg tccaaaataa tgtccaatan aaatgcgact catcactgtc aacgtaaaca 60
attaacaatc aacatctaataa ttcgggggcta gagactaaaa tagcgagatt acaanaaaat 120
ggaagactca attgataaat taaattacag gggaacaaaa ttgtgagcta naatggaaga 180
otcaatccga cggcctcagc taccgtttcg aactcattcc ggacaccaat 230

<210> 252
<211> 224
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 252

catctacaac agacctcctt aacctcagca gtatattcat ccacaacaga ataattatga 60
cctctccagc aacaggtaca atcccgagtg gagaatcatc ccaaccttag atggctgaat 120
ccttcacaac aacagcagca acaacaacaa ccttaatttc anaatgctgc tggcccaagc 180
agaccatacg ttctccacc aatccagcaa caacaacagc aaca 224

<210> 253
<211> 426
<212> DNA
<213> Glycine max

<400> 253
caggctccac cagttctagt gaacgtgctg cccataccat tggcgctttc aatgcggcgc 60
atagatgtga tcgcggcaat tgaacccaaa gcttccaatg ggcacgcta catcctagac 120
gctatcgatt acttcactaa gagggaggaa gccggttcat atgctaccgt gactagaaat 180
gtggcgggta ggatcataat aaaggagata atttgcaaat atgggctgcc gagctatatc 240
atcactaaca acgccgcaa cttgaataat aagatgacga atgagttgtg tgggtattcc 300
aagagaccac accataatct gactccttat cgaccaaga tgaatgatgg agttgacgcc 360
gctcacaaga atactaagat gatcatctta gagatgacag cgacatacaa aggattgcac 420
gagacg 426

<210> 254
<211> 505

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 254

ttgacccctg agntttgaga ccccgtagn atcgncgatn ctttggagaa gaantctagg 60
catgcaagct tttaatggaa gtatagagca tganagtgtt ctgataccat taactagtca 120
ataggggtctt gagcagggttg aggacatcca tactatattt ggaaagaccc aaaagaagga 180
aaaaaagagt aaaacttgca tatggaagaa gaggtcgata ttgtttgatc ttccatattg 240
gtttgatcta gatgtcagac attgtagcaa tgttatgcat gcggagaaaa atgtgtgtgg 300
tagtgtcatt ggcacacttc ttaacattca aggcatgaca aatgaggcgt agaacactcg 360
acacgatctt attaatatgt ggatctgaga ccacttactt cctaggtttg acggtgaaat 420
gttatectct cctcccacta gtgatacttt gttcaataac acattaactt tacttccatc 480
ttattctacg cttccttctc ctccc 505

<210> 255
<211> 432
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 255

gttgattact attttttaac atcgatttta gcgtcaaccg atataaaaag tgctttacaa 60
caccgatttt cattagaacc gatgtaaaaa gtgctttaca acatcaattt ttagtagaac 120
cgatattaat gtagtnttg cataattaaa aaaatatttt gtttcacaag aaccattttc 180
ttcgtgatcc atttttttaa aaacataatc ctgtgaccat gaagaacaaa actaagacac 240
tataatatta atataataaa tgcaacatga agcagctagc gaacttcaaa aaaattattg 300
cgaaccaatg atttgatgtt ccagtagcaa cttggaccct tggaatgctc gtattgctta 360
tagcatccta naagaataac aattattagt tgcattgggt gggttccaat tatccctgaa 420
aatatgtcag ct 432

<210> 256
<211> 206
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 256

gcataatcca taatgctgca naacannata cacgcatgaa taaactotta atcgacatat 60
ggagtacata ggaanatcct ttcttaatag ttccagcctc catagacaat aacgttgctt 120
atgaattaag aacaacaata aaactagtct gaatatgtga aacacataat ctgaagaaag 180
aatagctata ttcataca gcaata 206

<210> 257
<211> 221
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 257

tatagtaaca ggcanacgga aggattccac acatactaag ccttcattgg tgatgacgaa 60
tttgttgaat ntgatgagaa cctcattgca atcttcaagc acatattcaa cctttatgga 120
gaatattctg atagtgccaa aaagaagata gccagaact ctattgcana atcttgtctg 180
cagaagggaa aatganatg aataacataa cataacatta a 221

<210> 258
<211> 478
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 258

gcctcgccaa gcattcccat tgaagaacca ttattcanac ctttcanagt tagtgagaag 60
gctaaaagaa aaattaggga acttagaana actaaatcct taattgaagg agtaggcgat 120
aatcatggtg aattactaaa caaatnggta gtttacttaa ggatcatccca gaaactccnc 180
aaacttcaga taatacttcc aaaatggtaa caagaagtac ttncanataa ttaatgtatt 240
aatgaagaat gtgaccaacc tcaaaaatca agtggataga tcagttcaga aagaatataa 300
tcattaatcc aacactggaa acacctctaa tatatataac gccaaactgcc tgactttata 360
gagaagaggg aaacatntag agttagtga cacattatga tggactgatg caacgagata 420
catatgatcc tcacacatga ctgagtcact acaactcatg atgtaaaaca tatatatt 478

<210> 259
 <211> 507
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 259

agggatgact ccttgagacc atcgatcctc agagacgac tcgcaggcat gccagcttaa 60
 tgagtatgga agcgtcaciaa ataacttctt aggtttttat aatctaata ggaaccagaa 120
 tctcggaggg gtgaagacac ccactacctg cgatgagttc ccgggtggag tactctggct 180
 attgtaaagt gtatagtcac gatagcttcc attgttttagc actatcacgt tttacattat 240
 ggagaaggcg cagacagagc ttatttttatt gctgacgtgc acgcccaggc gcagcactct 300
 cgagagagtc ttttaacagt atacgcttct gcgccatctg tgctatgata attcgaggty 360
 aaccagaata ttcttatcat gctgctcaag aacaatgcta gagtttatat tctgagctat 420
 ctaanacgat gggacattcg ctactccgtg aaataacaac cgcccagaat tctgcacgat 480
 accttctcat ttattataat cccaaag 507

<210> 260
 <211> 290
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 260

gtttaagtct tctaaactgc ctatctatat tccatagcct ataacaactt ccgtgtgccc 60
 atcggcttgt gggtgacaag ccggtgacaa taacaattta atgcccact tgctccacaa 120
 agtctctcan anatggctta agaacttaga gtccctatca ctaacaatgc tccttggcat 180
 accatggagt ctcaaatct ccttgaataa cacatcagcc acatgggaag catcatcaac 240
 tttcttacat ggaatataat gagccattnt agaacaccta tcaacaacca 290

<210> 261
 <211> 428
 <212> DNA
 <213> Glycine max

 <400> 261

gattggtgaa tcttctgct tttattggtg accacagagt ggtacctgga gatatgtcgc 60
 ggcggtcacg agaccttggg gacgtcaggt ggggtgctat tgcccaaac caagcttgac 120
 caatcccgac ccaacccggg catagtcggt cagtgagaac ctgtgatgta cctaaacagg 180
 cgagctcctg gcagtcaaca gataaaagga acaaagacca caaagcaagg aggcttgtgg 240
 tagctggcca gctgtgaaac ttgattgata tgtgagatat ggtctctggt aatcgattac 300
 caagggtgga gaatcgatta caaggcttat aaatgaagac aggttgctaa catggtctct 360
 ggtaatccat taccatatgt tgtacccgcc aacaggctcg gacactgatt cgagaactat 420
 aggacccg 428

<210> 262
 <211> 421
 <212> DNA
 <213> Glycine max

<400> 262

taacacttta acgtgcatat gttagcaggg tgtaattaat tacacactgc ttccgtactg 60
 ctaaaccaga acaaaagatg attaatggta gatgtttttt attatcgata aatattaatt 120
 tattaatttt tgtaaanaat atagctcaca agagtaatat ccaattcata tatggttcta 180
 tattgtgatc ccaaaagatg cactgacattt ttttttcttc tttccacatg aactcatata 240
 tcactccta atacaatggg ataagtgaag attgtaccat acctcatata atactactga 300
 gatatgatca cgttcacaaa tgaatttcat tcccatttat tctctactct ggataacatc 360
 ctccaactat attatcaacg cccacctgcc cctgaccttt tttagaaaaa gaggtgtaac 420
 g 421

<210> 263
 <211> 493
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 263

ttgccctgac cctggacctc gagaacgtga tacttacact cacctgaaga tgcagcttga 60
 gaatatgtct atactggatt aacnattat agccttatca taaacgacta cacaattgct 120
 cttgagacaa cgactgattt attcaagaga ctgtacttta atcgattacc atgccatata 180

atcgattact tccctttgta tgcgtgtgtc agaagcgaac aagaacactt taattgatta 240
 ctttgagtat ctaagtgatt acatagtcct tatgttcttt ccaattttcg agaagaacgg 300
 tttaatcgat taccaagata atctaatacga tacatcattg aattgagcga ttaccttgaa 360
 gactcaactg ataacagacc ggtgtaggtg ttttctctat aaacaaccaa cttgtgctat 420
 gtataacaac acaacaattt gatctctagc agagcctgca tcacttggtg ttattaaata 480
 aagaaagaag cat 493

<210> 264
 <211> 509
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 264

ttgcccttgt anccttgntn acntccggag nncgctgat ggtatgcagt cgatctcggg 60
 gcatgnaga ctaagcgcatt agttggggca tttatcacta ttacctgtat cacatacctg 120
 atccgattat gacttgact gtaaagaatc acgagtgatt gatattatga tcctcgccac 180
 cttgttcttg tacgaacaca tcataaact tatagagntt gtactacaca cagcacactc 240
 tctgatttag ggataggtgc atgcttgaac tatgattgct gggattgctc gattgtagtt 300
 tgagctcttt tacgctgtaa actcgccgtg acttgagecc aacatcactt acatacactc 360
 ttctatatgc tcttactcgc ttcaaagtga atatgatttg gaggcgactt taactcgcgt 420
 gtgacttgcc catgacaact catatgactc ttctcaaact aatctacggc cctatgttat 480
 taccaatgtg agttgatatc atcctaagg 509

<210> 265
 <211> 485
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 265

atganccttg agccccctcg agngcccggg atccttagag tcgaccgcg gcatgcaagc 60
 tnggactcat gagcgggccg tttatgactt attgaattga gccttggtgt gttactagt 120
 ctattactta tggcaccgat gagagcgcac cgactactca ctcaacagcc tattgtaaat 180

<210> 268
 <211> 272
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 268

gctcttaact gcacaaggct cttaatatgt gaagagtatc cttgtgtaac cttcacncga 60
 cgaagacact gacanagact tatcttcttc ttattggaca cagtatggca ggctggcggc 120
 aagtaaatat tcatcccatc agaccttggga tgcaactgtg atcgtatgcc catatcagct 180
 agatcttaac gggatttcaa gccatccttc gtcttgccctt gaatggtaag gagcgtacca 240
 atcacactgt cacaaacatt gttctccaca tg 272

<210> 269
 <211> 451
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 269

cacgcctgtt catntgactt tgccaccaca agtgggtgac tctctatcgt actcacgctc 60
 caggacgcgt ctctgcaca gaatgcagaa ctctctcact gcgcaaaaagg agtgtataga 120
 cttttcttca tatatcgagg cccgcatgca tcttcgtgac actattgacc ataaagcatt 180
 cttgcacatg catgatatga aattctgac ctaatcatat gcctcctcan gttcatctcc 240
 aagtacagtt gacagtgtgt gtttcttatg cacccttaac tattgcaa atgatgcggg 300
 ccactataca tatcacgctc tctacaccag aggacctttt ctatcttgaa cacttcatat 360
 attgcgcca tgacctatcg catccccact attacgcctt gagtaccaca atgcttacca 420
 atactcctct ttctcattac tacacaaacc g 451

<210> 270
 <211> 262
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 270

gaactcttaa gagaaaatgc attagaatct cttttgatga atcttagann atattggacc 60

gtgataagtt cgcanaaaga gtcagaanag acacanaaca agtaaagtat ttatagatnt 120
 agcaganaga aatagatcca aacgattagt aatcactggg atttgattaa ttngatcata 180
 atacctttgt tntgcattnt caaaatcatg gtaatcgatt acaatatgtg gtaatcgatt 240
 atctcannat aacatagatc at 262

<210> 271
 <211> 146
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 271

cttcattttc tccatgtatt tcttcacatg tcttatgtga atgggtgtaac atgattctnt 60
 agaatttcca ccaatttaac ttgctataga agctagattn gatttcctct agttcanatt 120
 tcttggtctt gntcttgaac catgaa 146

<210> 272
 <211> 99
 <212> DNA
 <213> Glycine max
 <400> 272

cgtatcccg tggagagtgtg atcctttata tttgagagaa acagctatca tttagcattg 60
 atttttgcat gaatctctga agtatggact gaatgcatg 99

<210> 273
 <211> 227
 <212> DNA
 <213> Glycine max
 <400> 273

ctctctgacc tgggaatttc cgttcaactt attgaccatt agtttcgagt cgatgaacac 60
 gtatcttgta tttattatat tatttgtag cggaagcgct attgttaatg cttgttcaag 120
 taccaccca tatcaggatt cctataagat tctcgctcca catagtgttg tactggatgg 180
 gccattgaca aatgtaacac cctctaccct cacacataac gaataaa 227

<210> 274

<211> 418
 <212> DNA
 <213> Glycine max

<400> 274

accaattaaa gagttcaactt ccaaacatcc cagccttccg tattggtaca attgaaacat 60
 tgagaaataa tcaagacttg tgtggaaatg tctctggctt ggaaccatgc ccaaaaagcaa 120
 gtaaaaaatc tcaaaatcat aagactaaca aagtcataatt ggtatttttta cccggttggtt 180
 tgggtacttt aatattggca ttatttgctt ttggagtctc atatcgtctt tgtcgaagct 240
 cagagacaaa agaacaccag gatgcaaaac caccaggcca aaatctattt gtgatatgga 300
 gttttgatgg aaaaatgggtg tatgagaaca tagttgacac cacataagag ttcgacaata 360
 aacatctcat tggagttgga ggacaatgaa gtgtttacaa agaagaaatt gacatact 418

<210> 275
 <211> 508
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 275

atgaaccatg aancttgata cctctgagan ccggcgaact ctgagacgac ccgcggcatg 60
 caagcttggt gagatacctg gccaatctca ttttttctat ataaaggtaa tagaactaaa 120
 tcatataaat aaaaggcaga caaaatgtcc attgtaaagc atgaccacat caatttcatt 180
 tgcattagtc ttgaagacat aaagattcta catgttgcaa gaataagaat gtaacagtag 240
 aacatcccat ttctcagaga atgtttacct attccaaaat atggttgtct tcaatgctaa 300
 gcaatgtaga ccatgctcat tgtgcccatc catttgaatt tcaactcaagt catcaccaac 360
 gttaacaagt tctctagagc acctgactat cagacgacaa agatcttttg ataaagaaac 420
 ctctgacaaa gtaacaacat cagcttccac aagggtgaagg ttaatacttt tcctcaatga 480
 ctcatcatg catgatgcat acacaccg 508

<210> 276
 <211> 437
 <212> DNA
 <213> Glycine max

<400> 276

cttcaaacac ttgtgtaatc gattacgata aacctgtaat caattaaaac aaagagtttt 60
aactatagag gaaatcttct aacttttagaa cttttcttct aaccttaca tgatgatgca 120
tgatgcacat atgatatgat agagactaag atgcaacaca caatataaca atcaatacaa 180
atgccactca agagagttgg gcatgtaaaa aataaaacat cttaaagctc ttcttcaagc 240
ttcaaggcta acgtttcatg ttgctctctc tatctctaac aatattttca tggcacaaaa 300
catatatata tatatatata tatatatata tatatatata tatatatata tatatatata 360
tatatatata tatatatata tatatatata aaagtgaatg atatgttttt cacatagaag 420
gcgttccacc acaatag 437

<210> 277
<211> 519
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 277

cgtgttgann ccatcgatna actgacactc tataatactc aagctgggca gcanacgtat 60
gtatgattnt aagttggctt gggagtgggt gctgacattc ctttctagtt gcacctatta 120
ctaataattta tgtttgaatt cgcattgatt tattttgatt ttaacatgac ttatgtggga 180
gcagggatga tttgccccca ccttgatgat tcgtctttgc tccttgaggc cattctctcc 240
ttcggagtga tatggccgct cattgatcgt cgcaagggag attgggtccc taccaattta 300
gatgagagca tcatgagagc tttgtacggc gtcgaggtct ttctaacagc tgctctcacc 360
ctcgggtgatg gcttatacaa ctttgtcaag atttttagttt tctcaatcct tagcgtacat 420
gaaataacta agaaccgtgg aaatgggatg acgtacatgt caattcccac tctcttgtct 480
gatattagtc atgaaaatta cgtcacaatg ctcatccg 519

<210> 278
<211> 453
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 278

tatctccgac agccaatggg tgagtctcgt ccaggtagtc ccgaaaaaga ctggcctcac 60

agtgatcaga aatgagaagg aggagctgat tcctactcgg gtgcagaaca gttggagagt 120
ctgcattgac tataggaggg tgaaccaggt taccaaaaag gaccattttc cctgccatt 180
cattgaccag atgcttgaac gcctggcagg taaatccac tactgtttcc ttgatggttn 240
ttctggttat atgcaaatta ctattgctcc tgaggatcag gaaaagacca cattcacctg 300
cccttcggc acttttgctt ataggaggat gccttcggc ctgtgcaatg ccctggtac 360
cttcagcga tgcataata gtattttcag tggattttta aaaaattgca tagaggtgtt 420
tatggatgat ttcactgtat atggatcctc ttn 453

<210> 279
<211> 265
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 279

taacgatttc taattatgtg ggccattaag tctatcatat gctaacaata gccgagaagc 60
ccatgaatct ctteggnggc ggagtaggtg tctgccattg ccttggcctt ggctaacaag 120
cggngaagtt cttgactccc gttcaaggta agagcaaacc gatccatcaa catggttgcc 180
tcttagtgta aagatcgat cacccttctt ctagcctctn tttccgtgta tacttgagca 240
tactcatccg cgattctatg ctctg 265

<210> 280
<211> 456
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 280

naggtgtttt gatgataaca atgatgacaa caaatatga tgaaaaaaaa gctcaagtga 60
atcaaagaac atctcaagag aatcaagaac aagtcaagag ttcaagaatc aagaagaatt 120
caagattcaa gaagaaagcc tacaacaag aatcaagatt caagattcaa gatctcaaga 180
atcaagatca agattcaaga ctcaagattc aagaatgaag aaaagactca atcaagataa 240
gtattaaaaa gttttttcaa aactttgaat agcacatgag tttttgacaa aacctttacc 300
aaagagtttt tactctctgg taatcgatta ccatattgtt gtaatcaatt accagtagca 360

naatgagttt gaaaatgtnt tcaaactgaa tntacaacat tccaaatatt ttcaaaaggc 420

tggaatcgat tacaatgttg tggtaatcga ttaccg 456

<210> 281
<211> 264
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 281

gtatcacaat atatgtagct acttcccttg agaaaaaatt aatggaaaat taatttgatt 60

ntatatnnta cttcaatttt ttggcaatgg ccggattgaa aaatatttaa ttgaataagg 120

gtgttatatt gttgaatntg tcaataatac atgtgttgaa gttattttgg ttttttttac 180

ttgattactt agtacccttc tctatattaa cgtttattat anaataagtn gtttataact 240

atgactaath gttcatatta aaat 264

<210> 282
<211> 246
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 282

agcaattggc aatttatact ggaatattct aaattacaaa tgtntataag aaacaaacct 60

gcaaatgact agctacttgc tctcttatca aagcagggat attcatagct tcaacaatcc 120

tctttggctt tgcttctgaa gaataatcaa gggagaagat taattacaat tatcatagaa 180

ttaagccata tacatgttnt taaataccaa ctaagaggaa tacanatggt gaagaataat 240

catact 246

<210> 283
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 283

taaaaactat aaaatgcaag gtaaataaaa tgacattcag tttgtagata tgttgggtct 60

ttctaacaaa caagctgatg catagaaata tatttctcta atcaatcgtg ctcttgtgtt 120
ctatgttgta gcctaaatta ctaaaccctc gatccctcgt caggatgaat atccaagctt 180
tgtccgcaga tccctcattt aagactacac ctgatttaga cagccctctt aggtatagac 240
taacttaaac tgagtntcat cgcagatcc cttatgtaag actagactca cttcagtagc 300
ttaccaaagt taagcctatt taagccaaag ctttgaccgc atatccttat gtagactagg 360
ccaacctaac cagctttatg tacagcatat ttaaaccaac cttacctcgc aatccctcat 420
gaaggctaag ttaatcctgt tcatcaattc taggcag 457

<210> 284
<211> 429
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 284

gctcactaca agccttaggt gataaaccat gatattacca tacccttaan ggaatttggg 60
gctttggaat tgtnntggga ataagtgtgt ggggtctatg gttcatagga acacatgcct 120
tggtgactat gttcatgat gtattntggg ccatacttga tgtacattgc atattggcta 180
aatgttggaac atgctgaatg aaatgttggt tctcataggt aaaagaaaac aataaagaga 240
acagcaatac agttgagtga ataagatctt aaatggcaca agaagatga gactcttggg 300
tctactcttc atgtctaatt ctatcttgac tcttttattg cgtagtgttt taatatgcac 360
tatcccttng ctctctatcc tttggattac cactattcat attctcatac cttgccttgc 420
ccatacacc 429

<210> 285
<211> 544
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 285

cgactgattg atttgattca gtactcgnag acgctgatcc tttatcactc acctgcgcgc 60
atgcaagctg gcttctacaa ccctgtcaaa cttaagtgtg ctcgnagnng gactgacaac 120
ccacgtagag tctgaaaacg agagtcttca cagtgcacag tcatgactgt gatgagccta 180

gtcctactcg ggggcagatc atatggagag tctgctttga ctattaggat gctgcaccac 240
 gggtaccata aaaggaccat tttctcctgc cattcattga ccagatgctt gaccgcctgg 300
 cacgtcaatc tcaactactgt ctcccttgatg gtttctctgg ttatatgcag attactattg 360
 ctccctgagga tcagggatag accacattca cctgcccctt cggcactttt gcttataaga 420
 ggatgcccctt tcgctgtgac catgcccctg gtcctctcag ccagtcagta ttactatttt 480
 cagtgatctt atagaacatt gcatataggt gttatggatg atctctctgt tttcgatcct 540
 cttt 544

<210> 286
 <211> 451
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 286

acanaatcta tgtatccaaa acccctcaat ttaatggatt ntcaagtttt gagaagtga 60
 attgggaatg ggataaattt ggagcaaact ctcacctcac acaagtctat aacatcaatt 120
 taaacttggt caaactggat ttacacctaa aatttcactg aatcaaaatt tgactcctca 180
 accccaatt ttaccctaga aatggctctt tggtcagttt ggtcatttgt ttttctcttt 240
 agcacagccc anactttctc ataagtccta aatgacattt caagctagga ttaactcact 300
 ttaacctcca aataccacta aatccagatt tggccttcca actctaaaaa attcactctt 360
 tntccactca taacaccata atctcacctt ctaacccttg gtaattctac ccttatctct 420
 aacagtttcc ataacaattt caccacaaca t 451

<210> 287
 <211> 54
 <212> DNA
 <213> Glycine max
 <400> 287

gtgaaagggc tagtgatgtg cttgtgtgtg tgtgtgtatg tgtgtcttgg agag 54

<210> 288
 <211> 457
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 288

cccatcaact gccctaactc tttcagactg gtgattccta gtctcttgac cttgacttga 60
 tagaacctct ttttaagcga aggcgcctga ctcgatccca tgttttacia agtgaaacaa 120
 aaccaatgt gaatcaaaac tccgacatct atcatgggtg gaatggatga atgcttgaag 180
 aaatgcatat gacacagata ctttttatga atacgggagc ccgggaaatt gtcccccttct 240
 tagatacaac attttgggca gcatggcgcc tgacgtatgt atttaagaag gcgaaatgga 300
 ccctccgtcg gtttgacaaa gtgaggggac caagacacaa tccgtgcatg atgcatatgc 360
 ggaaggcaca aaacggtgat gtacatagta cgacaatatc cacaacaaaa tataagcaaa 420
 ggcatacatg acatttanga ctacatgcat gacagtg 457

<210> 289
 <211> 402
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 289

gctagagctt agctacacat acctctctaa tagctaagct cacctccttg agatgagaag 60
 ctagagctta gctacacacc cncataata gctaagctca ccncatgac ananaaaaca 120
 tganaataca aaanaaagtc cttactacaa agactactca naatgccccg aaatacaagg 180
 ctaaaaccct atactattag aatggccaaa atacaaggcc caaacgaaga anaaacctat 240
 tctaataattt acaaagataa gcggggtcatg cttagcccat gggctcgaaa tctaccctaa 300
 ggctcatgag aaccttangg ccttcctttg atctctagcc caatctactt ggagtcttct 360
 acccaatgcc cttgcaggat aggattgcat cacatgtcat ga 402

<210> 290
 <211> 457
 <212> DNA
 <213> Glycine max

<400> 290

cccatcacat gtgggtactag gtggcggtcg ggcgatgggtg cacaacaagt tttccacatc 60
 cacaagcgc gcataaaccc accatcccct gttgcccacc tccaactga, ctcacgtact 120

cccacgtagc ccatatcctc ttttctctca acaccgggtc cccatcaatc ctcccaagct 180
 ttcccaacat caaagtaaaa cgacattcaa acagcacaag ctatcacagc caagcaaaac 240
 agagcaaagg cagaaaactc tgccaaaaca ccaaccatat cacagctttt ctactttaa 300
 gactccaata acaattcctt cgttccggtt cattaaccgt tggatcgact cgaaaattgt 360
 actggaagtc tttagtacat aagcctacga tttgaccgtt gggatctact agcacacatc 420
 cagaactcat tgtacattac tctctccaca accagcg 457

<210> 291
 <211> 219
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 291

tattgtagcc gatgctcttt ctggcggtca tgcattactt tctatgcttg aaacaaaatt 60
 gattggtctt gaatgtttga aaagcatgta tgaaaatgat gaaactnttg gagaaatctt 120
 tagaaattgt gagaaatctt cagagnatgg tttctttaga catgaaggct ttcttttcan 180
 agaaaacaaa ttgtgtgtgc ctaaagtgtc tactagaaa 219

<210> 292
 <211> 544
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 292

ngggacgtgg atgaattcct gcattgtccg caccgggtga tcctntacag ccgacccgca 60
 cgcatgcaac ctttatgaac tcacggngga naagcctcga actttgacac tcccgacgcg 120
 tcacttatta gggtttgttt cgtggagcgt tttggcgacg atagagaggg cgtggaggcc 180
 gctctggagc tcctcggcga ggaggctgta gacgagacgg tgtcgtttga cgaggctctg 240
 gccctcgaac ttgggcgata cgatcttcac gttgaagtgg gtttctttgt cggaacttcc 300
 cttcacggcg gcgtggcccc cgtgctggta cgacacgtcg tccacctcca aaacggtggc 360
 ttccagcgcc gtttgaagct tcgagcgaat cctgctggct cgagatagca gcgcgctggc 420
 tcctctcgaa ctcatgggat cggatgaatg gtggcgctcg gttanggttt tgagaacgga 480

ggagtttccg cagatganaa catggacaga aattggggag ggaaagttat atttcccctc 540
tgcn 544

<210> 293
<211> 480
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 293

ntgaacccan agtaacaact atgacttggc tcaacaacac ttgttgcgct tactctcaac 60
cttcaaaagc aataactccc taattgatct tttaagattc cctatcctaa atgagttttg 120
tttggggaca aatacctaata acaaaaaaac tcaccaataa gtctctattg agaatatggt 180
tacatacatt ctatggttaa gtgaatttgg ctctaaaggc ctatagagtt atatatctac 240
aaaatgactt ctcttttccc caacatagtc atgagttntg ctactttaga aaagtaagcc 300
ttctttcttc acgtaagtgc agtttttttt tcttgtttct tagttgtatt gcttgataaa 360
gacttatgat gacaaattaa tctctcattt ccagaatgat tcatgtcttt tggagagctt 420
gtcacttcat atgcaaatat attgttgcac ggcattcctt tccttgcttc taatgtgtgg 480

<210> 294
<211> 424
<212> DNA
<213> Glycine max

<400> 294

gggagtttta catatatgac aaaggcgcaa cgcgttatgg ttgaaaatac ccttctggtc 60
tttgactaaa aatataggta ttagatctgg agtacagata atcaagctaa aggtataaaa 120
gatagcctat gtggatcata acactataaa ggtgtgacca ggctttacag atttctactg 180
ttattatatt ctgtcttttg ctctgactct gataattatc aagatccttt ttcatatgtc 240
tccgcaccgc ttcacattct aattcattta cgtgtatatt tctttacact ttagaaacta 300
catccatcaa ccatgccctt aacgtctaaa tctgtgacct gtcacgacac aagcagaagc 360
ggtccaacaa aagtgaaaag acacgatgag gactttccca tattggagag atattcctca 420
gacg 424

<210> 295
 <211> 353
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 295

cgacacataa ctcccaccgc atatagaata tcgggccttg tattgggttag ataccttata 60
 ctccccacaa gactcttgaa gaccgtggaa tctaccttct ctcccttcac aaactttgat 120
 agcttcaagc caccttccat atgtgtgttc acgggattgc aatcaagcat attatatatc 180
 ttcaacactt cttttgtgta gctttcttgt gagacacaga taccattctc ccgttggttca 240
 cttncattcc caagtatatg acatgagtcc atatttgcac atcaattcac agacatgact 300
 cttgagtctc aacaatttgg tattgcgata aataggcatc cctaaacaat aat 353

<210> 296
 <211> 436
 <212> DNA
 <213> Glycine max

<400> 296

ccagtttcct gaaaatatca aggaatctcg ttagatggcg gtcgttggtcc ttcttggaag 60
 gtaccacagg atatggtact ttgtatcct catttgaagc tttttcttgc ttcttctctc 120
 ttgctttctc acttctactc ttttctttcc cttctttatt tttttcaact ttttcttttt 180
 cttcattttc ttttcttttc tctacctcta tttcttttcc ttgggtggtt atttctttct 240
 tctcgaccgt tattgggtttt tcaactctct aactgtcac atctgttgcc tctcttttct 300
 ttttctcaat gccatccttt acaacaatat gtttctcaa agccacccta tctcctctct 360
 caactaccaa acgcttattt cttgtcatca caacattaca ttctctgtg ggattctctt 420
 ctgtgttcgc cccaag 436

<210> 297
 <211> 365
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 297

ctctacaatt gcatcacctc tcaatgatct ggtgaagaag aatgtggcat ttacctgtgg 60
 tgaaaaaaag gagcaagcct ttgcttttgc caaagaaaag cttactaagg ctncagttct 120
 agctcttctt gactnttcta anactttnga gctagaatgt gatgcctctg gagtgggagt 180
 tagagctgta ttngtacaag gtgggcactc tattgcttat tttagtgaan aacttcatag 240
 tgccaccctc aactacccca cctatgataa agagctntat gccttaataa gagccctcca 300
 nacttgnnga atataccttn gttccanggg aattgcattc atagtgatca tcaatcactt 360
 aagta 365

<210> 298
 <211> 453
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 298

aaggccaagt catcacgata tgcgaggatg actccccgag caagttggat ttggtatgac 60
 catgccctcc tggtttctga ctaggaaatt ggcgagtgga ggagcgccca gacattttacg 120
 cgacaagcat aatgtaaccc tttgtggctt ttaaactcta cggtggggcc taggcttttag 180
 agtttccttt tgttatggca ttatgtcttt tgttcttgaa tttataaata taaagatctt 240
 tcttcatctg ttcttgcacc tctaccatt ctcattcatt tgcattgtta tttctttacg 300
 cttaanacac tagatccaac aacgagtcct tcgaaggtag taatacctgn gacccggcca 360
 tcgattcaag caagaagcgg gtcaaacaga gagtgaagag gacgaggatg tgggacttcc 420
 cccagagttg gagaagatag tcactcacga ggt 453

<210> 299
 <211> 479
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 299

nggaatggca cttggaccac ctatttgaat ctctatgct gtacctacat acataaaaac 60
 agtccccacca tctcaatttt tacaaaatca tattcatata ccattggggc atttcaccaa 120
 gcacttgggtg agcgcatgtt tggacatgaa ttgcaagaga atgggagcaa tgtggcatgc 180

cccattgctt cagaatacaa cctaggccta agaccttttc attcaaatcc tcaattcaag 240
 aaaacaagca ccaaagcaaa ccaaaaactgc ctcacaaata taagcatgtt ctcacaattt 300
 aaggcaccaa aagatgaaga aaacacatca atgggaagca aaaacatcaa g gatggaata 360
 cttacttggt ggagtgaatt gaaacaccaa aaatgaaagc aaaaggcaac caatagtggc 420
 ttgagggggc aagaaccaca agccttcgtg ttctttcttt cttgaatgag aagggggan 479

<210> 300
 <211> 541
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 300

cggcacgatg tttgttgcatt tgcgtaccca gacctctag agtcacctgc ggcattgcaag 60
 ctngaacaga ttaaagtcaa caacactgac tgttntgggt gagaggtgga gacctgagtc 120
 acacatattt catcttctag ttggtgaatg catagtcact ttggaagatg ttgctcttca 180
 cctgagttta tgcgttgatg gaaaaccaat taatgaccta acatattatg attgggaaca 240
 aatgtgtgca aaatatatac gtgttggtcc cccaaagaat gcactagtgg gatcaacact 300
 aaaactaana tgggtaaaag aaaacatgct gactctccca gcanaatcca cgcaacaata 360
 attagcacc c attgtaggc atacattnta ggaccaatta gacaagtcag gaaacanagt 420
 tcacctgatg tatctacctc tgtagcaaaa tcttgaacag gcaggatggg acaattgaga 480
 attgacatgt ttagcacatt tgtacagaga aatgtttatg acaatttatc catcatcaaa 540
 n 541

<210> 301
 <211> 221
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 301

gcctttcttc ttctgaaca cacatgggtca ttaattcatt gatagaccac ttatctttat 60
 gtgtcgtgta ggaaatctta aatggcccat attcatgccg aagggtgttc agaataaat 120
 gcactangaa ggactcagac atatcaacct ctagtttctt aagtngagct gaaatatctc 180

gcattntcat gatgtactca cgcacacact tcacacttgt g

221

<210> 302
<211> 170
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 302

agcacaacta gtcctatgtc ttctctttga gagatanaga tactcataag agttgatgta 60

ctctactata catagttctc tcactgtgtn ggtcaacttg atgaactctc tcaagtgttt 120

cataggatct tcatgagcac caccactaaa tacgttgtct atagcatctg 170

<210> 303
<211> 256
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 303

gctcgaaaga gagttatgcn ctgtacactt anacagtgtt cagagtaata tattatgcca 60

nnaataagag aacatgacac ttggacccaa tgatgtccaa tntcacaaaa ctcaattaan 120

aggcttcana accataataa aacatgtcan atatatgcaa aatgaaacta taatgtatgc 180

tctaatatcc tctatcagag gacattcgat aggaacanaa tgaagtcctt tanacaatat 240

tcttattgat gatgat 256

<210> 304
<211> 260
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 304

catgtggact atgtggcggg cgggcatgg tgctcaacaa gttntccaca tccacaatgc 60

gcgcataaac ccaccatccc ctgatgccca cctccatctg agctcacgta ctaccacgta 120

gcccataatc ctggttgtct caacaccggg tgcccatcaa tctctgcaca gctccacaac 180

atccaagcga aacaacattc aaacagcaca agctatcaca gccaaagcaa acagagcaca 240

ggcagaaact ctgccaaaca 260

<210> 305
 <211> 523
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 305

 nccacgggna agttccccag nnantcantn anatnganan naancaaana naataagaga 60
 acaatgaaaa tggaagaatt gattcatgtt tcctttgatg agtctaattg tatttgtcca 120
 agaaaggata ttttagatga tattgtagaa tctttagaac aaatgcacat tcatggacaa 180
 gattctaaag gaaaaggaga aggaagcaat gaagatcctc cagtagaagt caaagaaaat 240
 aatgatcttc caagagagtg gaaagcttca agagatcatt cccttgacaa cattattggt 300
 aatatctcaa aagggataac aactagacac tctctcaaag atttatgcaa taacatgggt 360
 tttgtatcta taattgaacc taaaaattta aatgaagcca taatagatga aaattggata 420
 atagctatgc aggaagaact ataaccaatt gaaagaaata atggtntgga gttagttgag 480
 aaacctgaaa actaccaat cattggaaca aaatgggtgt tag 523

<210> 306
 <211> 468
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 306

 agactgagcg cttatcaciaa ggtctgtgct tagcggatag acaattgcaa aaaaaatttc 60
 taagtctttt tctgtctata tcttcacaca agcttaaaac cccttggtca ttactaaaca 120
 aactaaaatt aatcacaatc acaatcaaga tatcctaact acatgcaaga ggtaataatg 180
 aaaatagaaa agggaaagaa aagctaggtt gcctcccagt aagcgctctt ttaacgtcac 240
 tagcttgacg catcgctctg ttatccagga accaagagag ttctacttc aaggaccttc 300
 ttctcaggtc tcttttcttc catcacatgc actntanaat aaacattntg gctaggtgga 360
 tccttggtct cctgaaacaa atcaaagctg atcttctgat cttctatgcc catccgcagt 420
 atcttttttc ccatgtncac cacacagctt gcagtagaca tgaatggn 468

<210> 307
 <211> 472
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 307

aagctccttc aactgcacaa ggctcttaat atttgaagag tatecttggtg gaaccttcac 60
 ccaacgaaga cactgacaaa aacttatctt ctccttcttg gacaaagtat ggcaggctgg 120
 gggcaagtaa attttcttcc catcagacct tggatgcaac tgtgatcgta tacccatatac 180
 agctagatct atgtgtgtgg ctgtgtgtgt atggctgtgt gtgagtgtat ctgtgtatgt 240
 gtgagtgtgt gtgtgtgtgt gtgtggttgt gtgtgtgtgt gtgtgtgtgt gtgtggttgt 300
 gagtgtgtgt gttagtatga gtgtgagcgg ctgagtgtga gcatatgtgc atgaagcgag 360
 aaccatatct caacctctgt aattctatgt acatgcactc cccttcaact ttaatgcgca 420
 tctatcagca acctttcatt nctctccgta gaatgcttcg acaatccgcc cg 472

<210> 308
 <211> 475
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 308

cgattggtgc atcagtcctcg agaccctatn atcaaactga ggcggcgtct cactgagagg 60
 gcacgaccca gactgttctt attagagggtg gataccggac tcaccattac tataatctct 120
 acctgtgca tgcttggggg ctccagtaga tgccggtcat ccctgagct tatacgcta 180
 tggatatacca gatacggacc ctttatagca agagtgtctaa caaagcgtgc gatgcggata 240
 tacatgtaga tcgccctgat actgcctaata gggagcacgt tctggcctta tggataggaa 300
 agcgtgatgc cagctagtga cctcacgcac ttactagta gactttgtgg catgcatgtt 360
 agaccatcta cagtaggaac cccgcacgtg ggacctctc tggacatact ggacagcagc 420
 ggtcgactag attgcacgcc atccactgca gagaaggat aaattatcat atcag 475

<210> 309
 <211> 511
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 309

gtgatgatgc atcattatct acgngaacat actcaagctt gctcggcacg aggtacttac 60
ccgttgaaga tcgatgaacg ttgattatcg aatgaagaac gttgaagaac ggttgatacc 120
tttgagagat tcctcaccga caacgttgcg gatacgcatt cgaatcgcggt gcgcttagat 180
tgacttgatg tggacaagta atgcgagcaa attggattga cacatagggtg cctaattgggc 240
tcaacgcctt aattcttgtc tttctaactt atatataaca aaacaaggga cgtgggttgac 300
gccagctcg cccaggcgag ctcaactcgc ccaggcgagc aggggttgctt cctccagaag 360
caaccgcctt ctggaggaat attccggagg gcccaagtgt gcctgggtgc tatttgcacc 420
cccatcttta ctaagaacac cacgctacgc tgttttcggg gagggctctat aatacagtac 480
cgtaacttac gatcgtctga agaaaggggg g . 511

<210> 310
<211> 421
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 310

agttgcacat ggatttttct canaacattc ttaccaaaga gtttttactc tttggtaatc 60
gattaccaga ttattgtaat cgattaccag tagcaaaatg gatttgaaaa agttttcaaa 120
ttgaatttac aatgttccaa ttaatttcaa aaagctgtaa tcgattacaa tgttttggta 180
atcgattacc agttcctttg aacgttgaca ttcaaattca aatgtgaaga gtcacatcct 240
ttcacataaa agccttgtgt aatcgattac actgatttgg taatcgatta tcaatgatta 300
tttctgaata aatcaaaaga tgtaactctt catattgttt tgatttttca catggattaa 360
gctctctaaa actatactct tcatatgggc gatngaccaa cttaatagtc atacattttc 420
t 421

<210> 311
<211> 234
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 311

acatcatcta ctattgttca tctgcttcca tgaatgaaga ttcattgatca tcacaggtac 60
cacaccacag gtacaanaat tgcagggtga gtntattata aaagaaataa tcaagcatta 120
gatgacaata attagcaagg aaactataac aataaccata atcatactta ataattcatc 180
agtttgacat acactanaca tctagtcac aactttcatc atttncaatc aatc 234

<210> 312

<211> 221

<212> DNA

<213> Glycine max

<400> 312

aacagttcaa tcacatgcc ataaccacat cctgtgcccc tcaactgagct agactcacga 60
ccattctgtt ttaaacggtc catatctcca agttccacat aaggatcaca taccagccag 120
tatacagcag caacaggcaa gaaaacttgt caaaaccaca aataagtttt actaagaaca 180
gtacattctc atcaatcgta ccgtgatcga tcaaattatg a 221

<210> 313

<211> 455

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 313

ctgagcaaat tcaaacgaca ataactntat aatcggatgt cctattgagt cccctaatat 60
atcaaactgc tccaaattga aaatggaagc tcgtagcata tttaaacgag aataactctt 120
tactcaaattg tgcgattgag tcacgtaata tatcgagacg ctctaaattg aaaacggaag 180
ctcatagcaa atgtaaaccg taataacttt taactcggat gtccgaatga gtctctgtgat 240
atattgagac gctcataatt gaaaacagat gctctgcgca tattctaaca acaataacct 300
tttactctgt tgtgcgaatg agtactggaa tattgngaga ccctcgaatt gaacacaaaag 360
ctcctaaaaa atcaaacaaa aacttttatt ttatgttcac tgaaccgtat tttcggacgc 420
tcacatggaa caaacttctt tattcaacgc agtcg 455

<210> 314

<211> 446

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 314

tcaaggaagc ttcttaagga agtttctcaa ggaagctacc tagtctataa atagaagcat 60
gtgtaacact tgtggtaact ttgatgaata agagtcttgt gagacacaac tcaaagttca 120
acttctctcc cctttttcct ccttcaattt tgtgctcccc cctctctctt tcttttcctc 180
cattgaagaa tcctctccaa gcttcttatt caaggcacat tcttggtggc gaagctcctt 240
cttccatggc tntttcccta gaggatggcg cctcttctcc tttgtcttcc actgcatctc 300
cgtggtggaa aatcaccatt gaaggacctc attgaagctc aaagatccag cctncataga 360
agttcacaag taagcttcat catattctct tangcacaac actgtggcag tatggactac 420
cagcgacaat gcatcaccat naaaat 446

<210> 315
<211> 482
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 315

ctgatgggtg cgagaagaga tcacatgttt gtcacatca aaaaggggga gaatgtgaat 60
gtatgtatac atgattttga tgatgtcaaa gaagaatcta acaaggctac ttcaaagat 120
aagcatttgc ttcaagaata attcaagatt gcttcaaaa acaaagcctt gtttcaagat 180
tcactaaaga ccaagccttg ccttataaca aagtgtttc aagacatgca aggctctggt 240
aatcgattac caggaagtgt aatcgattac ccgaagcagg gttgagaaat agctgttgaa 300
aaaggttttg aatttgaatt ttcaacatgt aatcgattac catatgtctg taatcgatta 360
ccagcaacga aactntggaa attcaaattc aaaagtcata acccttcaaa ttataactgt 420
gtagtcgatt acacacacat tgtaatcgat taccagtggg gagtttcaga aaatctgcc 480
cg 482

<210> 316
<211> 441
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 316

attaggttgt cagttctgca aagataacaa tgcagacgta gcataattac attatagtaa 60
aaaaattggt ttgcagacac ataccttgat tttaaatttt tgaacataaa cgatcataag 120
ctctagaaaa aatactcacc agtctccaaa aacatgggtg gcaatggcac atggtgagtg 180
gccttggcct ccaatgtaat ctaaattagg gaaataaatc aaattctacc aaaagtgtt 240
ttcaaatttc aaaatgtaga cttaaaaaac acaaattaac actatgtcat catcctccaa 300
caaaaacaaa cctgtgtttg cattctatga agtgaactat ttgtttaaat gattatgcat 360
gaagacatta nacacagttg taccattaaa tatgcaaact atgagaatnt canacaactc 420
anataccaag ccaactattt a 441

<210> 317
<211> 262
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 317

cattctagca gntccttatg atataagcta agtcaatgac cagccttang ttttcgtatg 60
agggtgagagc atcagatcca acttcccttg atctacacaa ggatgtgatt aaagctggga 120
agcctanacg agaagagtta gatngagcca tcatgggtcat ttgtctagag atcaaaccgc 180
caatgttcat gtccatcctt gtgattaagc catagaccga cctagctcta tcntgtgtca 240
natctgaaat gaaggatgta tg 262

<210> 318
<211> 537
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 318

ggtgactcca tncnnnnatg aaacctctcg tagtaccgt gatectctag agacgacccc 60
gccgcatgca agctntgcgg atttgggtctt cgccagtga atgatcgaag tggatctgaa 120
aagaggcaaa tttaatcatc ctgcttagac gaatgagaaa actgnggcaa ataaagaggg 180

tgaggatgag ggagaaaccc atgctgtgac tgccattcct atacggccaa gtttcccacc 240
 aaaccaaca atgtcattac tcagtcaata acaaaccacc tccttaccca ccaccagtt 300
 atccacaaag gccatcccta aatcaaccac aaagcctgtc taccgcactt ccaatgacga 360
 agaccacctt tagcaaanac caaanaaaaa caccaaccaa gatatgaatt ttgcagcgaa 420
 nagectgtag gattcacccc aaattccggg gtcatatgct aacttgctcc catatctact 480
 tgataacgca atggtagcca taacccctgc tagggttcct caaacacctc atttttg 537

<210> 319
 <211> 512
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 319

nggcgtgcta tgtcctgagn atcnacngat ntgaaaaacc aagctttatt ttgcttgatt 60
 ctctgcaatt ctctgcatcc ttgctcctcg aattgaacct tcaacctttg acattatattt 120
 ctcttcttct actatgaggg aaggtaactt ctttgtgtga gtgcttttgg ttttggtttt 180
 gacgtaagta ggaggaaggt tataggtaaa aaaaaaatta taattatattt ttaatcattc 240
 tgtagagaag aagttgttta ttaagagagt gggtagagaga ggttaattaa taatggagaa 300
 gttataactc cctcctttat ctaattgatt cacatgatat tttagaacaa acacaatatt 360
 tccatttgta ataaacattc ataaaattaa attcttcgat taattagcat gaatggtagc 420
 agtatgaaga attattatatt ntatcaatga ggcaagagat agttctaatt ttataaaaca 480
 agttaaatat gctnttagtc cataataaat tg 512

<210> 320
 <211> 533
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 320

tggacaagtt gacgcttgta aactggacca tagagncacc ctgcggcatg caagcttatc 60
 gctgacttat tatccacaaa aagcttcact ccactactct ccttgattnt taattcctat 120
 aataatgtgt ccaaccagac agcttggcaa gcactcattg cagctgcaac atactcagct 180

tcacatgatg ataaagccac tatggattgc ttcttagaac tccatgatat tgggtgttgca 240
ccatacatga atatgtaacc tgtagtactc tttttgtcat ctctgtctcc tccccgatcc 300
gcatcagtat atcccactaa ttcttctgag ttgggtgttggt ctttatttgg aaatagaatt 360
ccagtattga tggttccttt tatgaacctt agaatcctct tagcagctag gagatgagga 420
attctggggtc ttttcgtata tctacttacc agtccaacaa caaattccaa atcaggtctt 480
gaatgataca agtacctgat gagagaacca acaatctggt tgaactcagt ttn 533

<210> 321
<211> 301
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 321

agagacatga aagaccggat gagntttact gtgagatgga agatctaact tataagcaac 60
aacaccacc ttatntaaca cctggaaagg accataaaat cgangggaga gnttttcatt 120
aatccttnta gccaaggatc ttctcttgta ggggtgcatct tcaagaacac ccaatcaccg 180
actgtgtatt ctatgtcctg acgaacgttt gtggcattng ctcgcatgat atcttgagac 240
ttcaacanga tttctcttat agtagccaat aattaatcta agcaannttg tagttattga 300
c 301

<210> 322
<211> 450
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 322

attatcatga aaccacccta naccaagaga acagagtaga ggcagaaaac tctgccaag 60
actcattcaa attccacagt tttccctact caaatacccc agtaacattc tcttagttcc 120
gattcgtaa ccattggatc accttgaaac gtttactgga ggttcctagt acataaatct 180
acattttgac cgttgggatc tactagaaaa tatctagaac acgagatata ctacctttcc 240
cgtgactgggt gctgcacaag cattttttct gcacatttgg tcaagtttgc tgcacaattt 300
gacagctttt gctgcacaat ttggcagatt tcanaatcca actttcccac antccaattt 360

actcannatg gatcctanaa ttcctaaaatc atgtataaat catanttaaa ccanaaacia 420
 acttcagacc aaggcanatc anaatatagc 450

<210> 323
 <211> 362
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 323

ctcagtatgt tccttgggta gataattaat aattaactta gatacaattc acaattgtat 60
 ataancatta natatgttat aattaaagaa ataactacct ctcttgccca cactttggct 120
 accacatgat taacatatga tgtcaacact aatgtatctt ggggccacc tggaaaaccc 180
 tatgaatcaa cacctacatc ctttgaatt ggatcatgan ngttctcatg agtctcatca 240
 gcagcatcat cgatatgccc attatcctcg acaatagggtg cagttgttca ttatctacgt 300
 gtcgacactt tcaatcttcg acgctgangg gcttcttctc gtcaccacta acctctctac 360
 ct 362

<210> 324
 <211> 531
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 324

tgatgaatca tctcganctt gagatcctta gagacaccct gcggcatgca agctntgacc 60
 gcactttaac aatatctttt gttctattnt tgtgttgttn ttaatcaatg tctttaagtg 120
 catcttaaca atgtcttttg gatagggatg acaaaataga cactaatttt gtgggtatct 180
 ataaaagtat ttgcaaatag gaaggataat tatctgctta ttgggactag agatggggcg 240
 gngatactat agtaccatct caccctccc cgcacatgta tgatcatatat tttatatatt 300
 aatgtaatta aaaaataatt ataatttctt aattttatga ctagcaataa caatctaaca 360
 aagaataaag aatcctaatt caacattgtt atattaaatt tgcttcacat tctacataag 420
 aatatcgaac tactatcttac tgtttatgta taaagtaatg actctgctaa caagttatta 480
 taagattgac ttgtgaaatg gtattctgtg ttgttcgatg cttcaaacta g 531

<210> 325
 <211> 470
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 325

atttcaatgc ggaaagtatt atgttcttca ctatccatgt tcacacatta ttgctgcttg 60
 tgggttacgtg aacatgaatt acttccaata tgtagatgtt gtttacacaa atgagcacat 120
 cctanaagct tattccgcgc aatggtggcc tcttgngaatt gaagcgagta ttctctcttc 180
 tgatgagcaa tggacactta tccctgatcc aagtacaatt cgtgcgaaag gtcggccaaa 240
 atcaacaagg ataatgaatg agatggattg gctggaccat ctgacaccga caanatgtnt 300
 agatgtgaag agaagaccac agacgtgatg tcaatgaatc tgatgtggaa gttgtaataa 360
 tgattatgta tttgttgtca cttaatgaat gacctatcat gacagctgtt ttaaatagta 420
 tatatattat ggcgcctaa ctgacaatgg taatatatac ataatgatat 470

<210> 326
 <211> 315
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 326

ctgcagctag ctgctggcag ctggtggaag ctccttctct atttnnncta taatagggga 60
 ggagtgaagg agagaaatgt tcagaccttc tgggtatttcg agatcacttg aaattagtga 120
 aaaanactgt ctccgtgaag aaaatacaag ccgacgcgct ttcgtaacgt ttcgtgggga 180
 tttcgcgaag aatttaccta tntcttcgac gtcttcgttc gttcttcggt cttcaaccgc 240
 gtaagttctc gaaatcgaaa ctttcaattc attctatgta cccttagtgg tcttcatttg 300
 tttcacgtgc tttat 315

<210> 327
 <211> 352
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 327

ctctctctnn tcgaattggt gaggaagatt atttccgtga agaanatcca agccgagggc 60
 gcttcgtaac gtttccgtga gtaattacgc gaatattctc gaccgttctt canagattca 120
 tcgttcgttc ttcgttttct tcagtcttca acgggtaagt acctcanaca aagctnttca 180
 attcactcta tgtaccctgt gnggtccaca ttntgggtca tggagtttta ttctcattnt 240
 catttactnt tntatacccc ttttgacgtg cttaagccat ntatttaagt catttctcgc 300
 ttaatctacn aaataaaaata aatttccacc gatcggttga atcgcatatc cg 352

<210> 328
 <211> 514
 <212> DNA
 <213> Glycine max

<400> 328

ctttgactgc tgcactctgc acccttagag acgaatcgag gcatgcaagc ttagcaccac 60
 tatcgcgctt agcgcgagta aatggatttg gacttggcgc caatgttgca ctgagcctag 120
 caagagatgg acgactcgt tagcgagctg atctcgcgct tagggcgctg cttcgattca 180
 ggtgctcttc cagattcctt tttcacgcta agtgcactga agccgtgctt agtgacggat 240
 acgcactaag tccactgagt tcgcttagtg cgacaccag cttccgactt gaagacatca 300
 gtaacttatt atcttagctc ggccaaagtc tacctctctt catctcacag aggccccacg 360
 catcgtacta gcaccgctgc ctgttattcg tacaagtagc tgacaactat acacaggtac 420
 ccttcacta tcgcatctac ctcaacgcaa gcatcagcta ctgtacgtat cgtctcatct 480
 ccgcatectc tcgcgcactg ccacgagacg tccg 514

<210> 329
 <211> 341
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 329

tctacacctg ttgcaagagt ctgtggtcta tgttcttcta cagatcacca tacagatctc 60
 ggtccttctt tgtagcaatc tggagtcaat gagcaacctg aagcttatgc tgcatacatt 120
 tataatagac ctctcagca gcaaaaacaa caacagaaaa ataattatga cttttcaagc 180

aatagataca atctaggttg gaggaatcat ccaaactga gatggacaag tccttcacaa 240
 caacaacagc ttatcgcttc tttctagaat gctgctggc caagcaagcc atatgtntct 300
 tctncaatac agcaacaaca gtcacaaana agacaacaag c 341

<210> 330
 <211> 450
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 330

atgtatccaa aacccctcaa tttaatggat tttcaagttt tgagaagtga aattgggaat 60
 gggataaatt tggagcaaac tctcacctca cacaagtcta taacatcaat ttaaacttga 120
 tcaaactgga tttacaccta aaatttcact gaatcagaat gtgactactc aacccccaat 180
 tttaccctag aaatggctct ttgttcagtt aggtcatttg tttttctctt tagcacagcc 240
 cagactttct cataagtcct atatgacatt gcaagctagg attaactgac tgtaacctcc 300
 acataccact aaatccagat ttggccttcc aactatanaa cattcactct ttttacactc 360
 ataacaccat aatctcacct tctaaccctt ggtaattct acacttcac ctaaacagat 420
 ctccataagc aagttcagca cacatacatn 450

<210> 331
 <211> 471
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 331

cgtcagagtt tanggattga gcttggttca actgagtgc catctgcccc atctgattgg 60
 tcaaactctg aatggaggct cttgtctctt gctgaaactg catattttgt attgtcattt 120
 gccttactaa ctctcttaag gaaggttgag aaggggcctc agttgattgt tgtctctggt 180
 gttgttgctg ttgatgctat tggttggtt gcattggagg aggaatgtat ggcttgcttg 240
 gaccagcaac attttggaag gaaggagcag gttgttggtg ctgttggtga gggctagacc 300
 atatgagatt agggtgattc ctccatccga gattgcattt gttgctagag aggtcataat 360
 tgttctgcta tggctgattc tgctgctgag gatgaggag tctattgtaa atgtttgcag 420

catatagctt angctgctca attgctccag attgctgcat agaanggcac g

471

<210> 332
<211> 446
<212> DNA
<213> Glycine max

<400> 332

gtttccgttg ttcaatttcg agcgtgtaga tgagttatgt ccccgaaatcg gacatctgtg 60
tgaaaagtta tgaccattcg attttctcga gagcttccgt tgttcaattt cgagcgtctc 120
gatataattat gaccccgaaat cggacatctg tgtgaaaacg tatgaccatt cgattttctc 180
gagagcttcc gttgatcaat ttcgagcgtc tagatgagtt atgtccccga atcgaacatt 240
cgagtgaaaa cttatgacca ttcgaatttc tcgagagctt ccgttggttca atttcgagcg 300
tctcgatata taatgtcccc gaatcggaca tccgagcgaa atgttatgac cattcgatct 360
tctcgagagc ttccgttgtc aatttcgagc gtctcgatat attatgtccg cgactcggac 420
atccgtgtga aaacttatga ccattg 446

<210> 333
<211> 400
<212> DNA
<213> Glycine max

<400> 333

gagtgattca agaacaccct gtctgtatca tatgacattc acaacctttg cgtgttgccc 60
tcgctggaaa gagcgagtct ttcccttctt tcatctatac ccgttgatct ttcaaaccac 120
aagtcagaa gatacacctc tgcccagaat tatatcgtgg ccataactcc cattctacgc 180
actcacatta agtgattctt gagcctatac tgaatttcac aacgagttct ttcacctcag 240
tatggaacac ctcatggag ccttagctca gtatgtcatt ctaattttgt caccacactt 300
actagtttac atccattatc atttatgcaa gaccacttat agacacgaat acactattca 360
ccttctataa tccctttcat agttatcaac atctagcact 400

<210> 334
<211> 457
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 334

tatacaatgt tntcatgata aaagcctctg ttgaccatcg gaacaagatt tccaaaggg 60
gcctctggaa atgcagtcct tgcattcatg aacctgaact gtgagatgaa gatcgaagat 120
caaagatcgg atgggttcctg tctcgaatac ttaatttaaa ttagattggg ggagataaaa 180
tataattggg ctgatcttca tacaatgggt gcgattagcc aaatgcctga atatgggaaa 240
attccctgac tgccgtgaat cccaatccct ctaattaaat aagctttaga atgcagcact 300
gaactaaaaa atttcttgcy tcatttatat acaatctagt aagtaatgca ccaaagttcc 360
aatttttgca tcanaggaca gagctgatag cacataaact aaatggcata aaaatacaac 420
caaatcttat cactgagttc tatcaatgga ggaggcg 457

<210> 335
<211> 260
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 335

ctatcttaaa gtaagctcta tgaaatttat aaatcgatat ttatatttct aacacgatct 60
tccttacaac tgataaagac aaagagaaca aattcaatta taaaataaag gataaaggaa 120
aaaactcttg acccacacac acaaaagaga cgtattatag taaaaaatat ttaaggcatg 180
aaggataagc aagagtggnt ntaatttata attctgacgt tctccttgta attcccattc 240
ataattttct tctgtacttg 260

<210> 336
<211> 446
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 336

tccttcaa at aacttgcaaa ttatgggttct agaattctat aattttctat aggtttcagt 60
ctgctagcta gttattagac tggttgattc tgaatagggt ttatcccaga tttattgtat 120
tttttgtttc taggtaatgt ccagtcatgc ttctttaaaa tctaattatt tgaaaccaca 180
aaatttagag cctgcacttt ttaatgtgtt tatgaaattc ttcgttttta gctattcatt 240

atccttgcaa tctcttattt cacatggtag tacaattcan attaacttca gattttgggt 300
 ttattttgat gtttgccttc gaagcagtct ggaattttcg gtcctttag acgggttcgg 360
 ntaatccttt cttatnggtt tgtctgtact ttgttagaag cactcaagac tagtgtggac 420
 gcttcagtcg ttcccacatt tgagag 446

<210> 337
 <211> 255
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 337

gatctgtgct tgcaactgct ntaagtagtt aatcatctca tctaacattg aagctntatc 60
 tgtctgcaat ttgccaagt aatccaccac gtacaggaat aatcaatcca catatataga 120
 acnntgattg tgcattngat tntggatnga atatgnttnt aactatgtaa ttnttgataa 180
 aaatgaaatg atattagtct ccgattctaa ctttaataca tttaatcctt annatttaaa 240
 agactgttnt cgtca 255

<210> 338
 <211> 441
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 338

ganaactacc aaaactaccc atcatatctc caaaacccc ataccacga aatttaagag 60
 agaaagaagt ccacccaaac ctgaaatttc gaagtccac tctagccac gcacttcacg 120
 actccaaaaa tgccctcctt tgcgattttg gagcagaaat gagcaccaa gggtgaagct 180
 ttgtttggag cttcaatgga gaatgagga gaaagaaagg caacgtgagg aagagagaga 240
 gctgtctgaa aaaagtgtgg gggctgagtg aagagagaga anagctttnt gggtataaaa 300
 taaaagggtt ttctcttttt ctattattnt attcanactc tgccacgtgt ccctaattga 360
 gtggagcana agggcccact ttctctttta ctgtgacca cactcagcca canaagtgag 420
 aanaatctga cctttganac t 441

<210> 339
 <211> 535
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 339

agagccgata ggttgggngc ttgnacnant cgacantata tagtactcat tgctgccaag 60
 aaggtggtgg atctcgaggc ccgactgaat gaattatagt ccatgctcaa ggagtctgag 120
 ctacgggctg ctagagagag ggaggccagc aaggagcttg aggaggagt tttcattttc 180
 aagaaggagg ccgtggagca gcatgaaaaa gggccttaac aagccgttgg gcaggctggg 240
 ttcttcacca aggaccttga cttgggtctc ttgaccctt ttaaggacgt gaagaatggt 300
 gttttgcttg acaaagacga tattgctgct aaagaggagt aaggcgatga tgccattggt 360
 tagggtgcct ttcgtttatt ttcttctttt ctccattggt tgaatttagc cgcattgggccc 420
 ttgtaattat gacaaattat cttcataage ttctcttctga tgacaaattn tgactatnt 480
 atgtatgtct tgtgtgttgg ccttatctat gcaatgctcc atgcttgtgg tagtn 535

<210> 340
 <211> 406
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 340

tacacctata ctcgccaactc ctttcacagn caantaacac aaacaanagc cttcctctac 60
 tacctgtgtt ngtgcagacc ttacaatcac tgccacanat ccattttcat gagcaacgga 120
 tcgagccac cgcanaacat nctctcggt gtcaatacct acaaccaatc agataatatt 180
 agttctacaa catattattt atttataaat actccaacat aacgtattac taaaagattg 240
 acgcatcaac aacaatggat tatcacccaa ttttctattt ataataatc acttttaaca 300
 taacttatca tcatgactcg catatacata ataaaattaa ccataaacac ccatcaacta 360
 tctacatcaa cttctatact ctctgtaaca gttataaaca ataata 406

<210> 341
 <211> 466
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 341

ttctacagaa tccgtanagt ttcgtgaggt ttcggaagga aaacaaccaa acaacacaaa 60
aattggggggg ggggggtgaac ttatcaagct aggagtgtaa atagcaattt tcaaattcttg 120
gcagaggatt ctggaccttt tctttcttcc tggctaagca acttgggcga gcagggtggc 180
aagcacctcc ctcattttgt tgaaaaatgg cttccggcgc ttccgtagaa ttcccgtaac 240
cataaataag tatatttcac ttaatatggg tgagaaggaa gaaaaaaaaa gaagaaaatc 300
aagtccgata tgcttccgta actttttcgt aaattacgaa gaaggggggt gaacttatca 360
agtgcgaggt gtaaatagaa atttttgaac tttcgaatct cggcccttcc agaacattat 420
ggaagctcgg gttgcttagg agggagcagc ctacctcgct tgggcc 466

<210> 342
<211> 444
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 342

cataacattt ctaagtgata aagttttctt cttgacttct ctttgggttc tgetcatggt 60
ttcttttaca agtctagttc taggctgata tgcagtggag ttggaaatct ttattccagt 120
aatgaccttg ggtatttttt ctgttttctt tcttactttg nttgtgagga tatcttattc 180
tcctctatac tggacacact tcctttctct cttcattaaa ctttcttctt tatagagatc 240
tgaacttgty gaatttgtyt ccagggtttg caacagttgc agtctatgac tctgatgacc 300
aatctcataa ttttctttgt gtttctcttt tggattttct caattacaca tcgcgttggt 360
atactctggt gattttaatc ccttctatac tagttgntca ttgtgcttaa tatagcttct 420
tctattttta tctcatgcag cttg 444

<210> 343
<211> 488
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 343

atctccttct tcactacatc aagaatcacc gngttgagtc ttctctgtgg ctgtcttact 60
 gggtagctc catcttctan atntattcga tgcatacatg tggatgggct aataccagga 120
 atgtccgcca gggccagcc tatagccttc ttatgcttct tgagaactga caacaacttc 180
 tcctcttgct catcagcaag ggaggcagat ataactactg gagaactctt gctatcatcc 240
 aagtaagcgt attntaaatn tgatggcaga ggcttcaatt ctgggtgtggc eggctggaca 300
 gtggtagaag gagatgggtt ctcagccttt acctcataaa gaaagtcaga ggtatgtgta 360
 cttctgaaa catgggttagt cctatctgac tctatnaaat caatctcaag aggtanaaca 420
 ccaccaccag gcatgcantc aatatcactc tcagaatcac tctcagcatc anattcagac 480
 atatgatc 488

<210> 344
 <211> 532
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 344

ntttaatccc tcagtgcccg agctccttaa gaaacccac gcatgcaagc ttcacgggtga 60
 atctacagtg attcanagat gttntgatga taacaatgat gataacaaaa gatgatgaca 120
 aaggatgatga caaaaagctc aaagggtcaat taaaggatga gttcaagatg ttcaagaagg 180
 aatcaagaac aattcaagac tcaagacgaa aagggtgaag gacacttcaa gattcaagag 240
 gaaagttgaa ttcaaaaatc aagattcaag gatcaagatt taagaatcaa gatcaagatt 300
 caagattcaa ggttcaagaa ctcaagagaa gacttaatca agataagtat ganaagggtt 360
 tttcaaaaac tgagtagcac atggattntt cacanaacat gtttagcana gagttnttac 420
 tctctggtaa tcgattacca gattgctgta atcgattact agtagcaaaa tgttnttgaa 480
 gttntcanat tgaatntaca acgttcatt taatttcana aagctgtaat cg 532

<210> 345
 <211> 290
 <212> DNA
 <213> Glycine max
 <400> 345

catttgcgty cttatttctg tatggatga gatgaaatgc aaagggttacg acttgtgtta 60

gtcgggttata atggaatgag cctaaacact tgagcttgag tgaaacgacg actgtgagggc 120
 tgtgggttgag gatccttcct tgatatctgt cattctcact agcttatttc aattatgact 180
 ctaatgcata tctttctatc tttgaaaagt tgcattgatg tgagaagcaa ttgattgaag 240
 cattccatga tattcatttc atatgattga atttttctgt aaacaaacac 290

<210> 346
 <211> 377
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 346

tatcacatat atatatatgt tatgcgtaca acatatatca ttacgcaatg acatttgagt 60
 ataataaaaa atagttctgc agggcctaac atttcagtgc ttatattaat ttaggtacca 120
 cttaacattt attattgagt caactctcta acgnatattc ataatttctc tttgtaatat 180
 taatttaatt ggntaaagaa acatatttct tatggataat aatggctttc agnttcttag 240
 tgaaccacat ctgannaata tacttgacaa gaaatgtgtt actatgtcat agntaatctt 300
 tttttctttt aaatacatca tctcttatgg acgatttttag actcggagga cttattatat 360
 ggacatacac ttatata 377

<210> 347
 <211> 396
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 347

cacacatgta gtgaacatct atgataacct gcttcgtgta agcatttgtt tagctataat 60
 ntatgaagaa ccacttctag ttctataatt gtacaacata ttagcatatg ccanactatg 120
 tgtatcattt ggatcaccag aataagaata ttacctcaat aaaatctcct tttggcatta 180
 gtgctctgca tgcattctcta ntccttttgt atggatgat taaactagt atgcaaataa 240
 caccagcatc tgcaaagaag ttagccacct cacctgaana tntaaatcgt gatgtctaan 300
 tattaataaa acaataaaat ataatcggaa gatatcaggg anagcattta gaaagcaaca 360
 taagaaaaaa cagaataact caccaatcct tctaatt 396

<210> 348
 <211> 225
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 348

 tagttacctt cttatgccta gccctatana tactcanaaa ctcttaattn taggagaatt 60
 ttgtagaatt gaaattaagn tgtgcttaga gagagcatta gcctcttctt tggtnnttgac 120
 tagaaaccaa atggattctt ctcaaagaag ctattccttt atggcaaata ctcctactcg 180
 gtatcgattc ttcattggatt gtggcatcgn tctgtcatct tctca 225

<210> 349
 <211> 203
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 349

 acgttatcta tagaacattt ccattggatg taatggatga aattgtgcat ctttaggtga 60
 gaaagaggct atgttttgaa ttgcanaatg tagcagttgg gctaaacgca tatccaccgg 120
 taagcgcaat ttcagcgtgc ttagtgcana ggagaatctg ggagagcatc aacatcaaag 180
 ccgcgcgcta agagtgggat tag 203

<210> 350
 <211> 455
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 350

 gacctaaatg aagactaaac atgcattggt tatgtaattg tattcattat gcgatataat 60
 ttgttgtaag ccattaataa ccaattaata ttattaagta ctggttttgt taagcaaaaa 120
 aattgttggc ccaacaaaaa tcatttacgc gtgtaggata catcattgtc ataattgaca 180
 acacataatg acatgcatgc gtattaaagt ttgagcgcga caccacattg actaacttga 240
 ctacacattc tgaaggaaac ataaacacga aacatgttca tgcgtgtcta atttttttga 300

aacaaagaga agcaatctgt ctgtgacaac catgtatata tatagcagac acaactaata 360
aatcacacat tatcttgctn tcacatagtc tcccaatgga tacacanagt atgaaatttg 420
tagagaaact agcagtcaga tgattgcaac tcacg 455

<210> 351
<211> 483
<212> DNA
<213> Glycine max

<400> 351

atgttagtct gctcacatca aagagatata ttgtcttctc tctcagatat atttgatcct 60
aatcttatcg ttttcttata tgcgaactca tcagctgtaa cattcttata ttatctacac 120
acttgagggt atatttatag taattaatac aacttatatc ttatctttga ttaacctgtg 180
cagattgtta agctttcgag aattaaaaaa agaattagac tcttgaggat cctgaattaa 240
acgtgaacaa gtatatatag gagcaagttt atccattgta atcattagat tagaaataaa 300
aactacttgt ttgtgataga aataaatatt tatagttaac ccacaatgaa tttcgggaata 360
ttattattga taattttata gtgcaaaggt atttacatta tatacactta ggctcataat 420
tgtttgccca gaataaagca acaaactatt ccaactataa agggaaataa gtcagaaata 480
aat 483

<210> 352
<211> 448
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 352

tagcattaaa tgcacatcct tcttgatgca acaaaattac tctgattggc ttttgtgtag 60
aatacttcag tataaaacca cttttccttg gtcaaagcat gtttttcaca tctgaatgag 120
tcttttgatg ttgttttaac aattttttta ctttagtttc atttattatt cataggattc 180
gacaaatcat atgagaatgt ctctccaaca tgaatctcag acacagaaaa ataaatatag 240
agcgaaatat cattttttta tggtgtatca ggatcatgact tggctctatc ttcattctaat 300
acttttgacg catgatgtat acaacatgat ctgatatcgc ataagatata actattcacc 360
ctcgtattta tgtgcatcga ataagaacaa ttatgagtat tgatttatct tatgacataa 420

atgtcnttat cttaacatag atgaaatg

448

<210> 353
<211> 277
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 353

cctggagata tgtncgcggg gtcaggagaa ccttgggacg tcaagtggng tgctatngcc 60
canaaccaag cttgaccaat cccgacccaa cccgcgcata gtcggtcagt gagaacctgt 120
gatgtaccta agcaggcgag ctcttggcag tcaacagata anaggaaaac aagaccacan 180
agcaaggagg cttgtggtgg ctggccagct gtgaattttg tgtaatatgt ggattgtggc 240
ctcttggaat cgattaccaa gggtagtaaa tcgatta 277

<210> 354
<211> 302
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 354

cggtgatact gggtttctaga agtgggtcca gccattccca tgnnttcaca tttaaagaaa 60
agggagaaac aaccaacctt atgtactcct tgctaattgtt gggttnttta acaatgtcac 120
acaacctcaa gaatctcctc anatgcatca caagaagctc tgnntgaact ggtactntta 180
atcatctgga ggaacccatg tgtgaactca naagtaactc ctcgaggaag tgggtgtagt 240
tcaatgggtg tggttttctcc caccagatgg gtgacataat acagcaatgt cagtgtagca 300
tc 302

<210> 355
<211> 436
<212> DNA
<213> Glycine max

<400> 355

agatgaacaa ccaaatgaaa catgacagtg aagaataaag gaggaaatat catttccatg 60
tggtataaag tgagaacaac ttgattttgt aattagccta aggtcttaac ttccaataat 120

taagccacct atattctatt ctgaatgact actactcacc aattatctgt acgccccccc 180
 ctcccatccg ttcacggata gcacacttgc gtgattcggt gttatatttt actcagcggt 240
 cctccgcgca cccctcacgt atgcatactt gatacccttc ggatcgtctt atacaccctc 300
 cgtccttgcc tctccgtctg actctttcta ccgactatgt atcgtgatga ctcgtattcc 360
 ggcgcgttct cgctcacgag ccactctgct atctttgagt ctcttatctt acttagctct 420
 acgtctttat agctct 436

<210> 356
 <211> 350
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 356

cttctagttt cactgatact tgtttactat tactggacaa catatctagt tctatgtctn 60
 tcannattgn ttattaaaaa gctttcattt gtganaaagt tattatcttt gattaatgca 120
 ctattcaacc cttcattcta gtgtgatnnt tggatattca tcatctattt taaaatcgag 180
 acatttgatc attctaattt aaaaattctg caatcttggg ctcctattt cataatacaa 240
 acatntgggc tccatatnta agagaatctg caattctggg ccctatattn tagaaaatcc 300
 tcaatcttgg cttaatcttt aattgtgtct acattcttat ttcttacttt 350

<210> 357
 <211> 505
 <212> DNA
 <213> Glycine max

<400> 357

cttgaacccc atctagtacc cgggacctc taagccacct gaggcacgca agcttgaaca 60
 ttctctcatt tgattaagat attattactc tacattttta gactgagact tatgagaaaa 120
 aacacaaaag ctggtgggct agactattag cctaggacaa gataaggctt gaagggggcc 180
 aagttttatt tgctcaatcg aaaatgcgaa ctaacaccaa tccaatccgg ttatacttat 240
 tatgtcaatg aaatcactat taaatcatct aaagtcaatg agatatcgta tgaattgttg 300
 ctattaacta acacatacac caaagactag aacaacgaat tgatttagca tcgaatatga 360

agagtgagga gcaccaacaa cattggcagt gtggcataat tttctgcaga cgcagcatcc 420
 accattcgtg tctttcgttt gatggtacat tgctgatgac gccccagag aaaagaaagg 480
 attggattga atgaccatca ttgct 505

<210> 358
 <211> 460
 <212> DNA
 <213> Glycine max

<400> 358

caacacctat tacaactctt aatgcatgct ttcttggtgc atttaagcta tcacccccaa 60
 caaatttgct taaaccatat gaattcaaac tcgtggtgag ttaaccctct tttggaccat 120
 caacgttgac ctcaaccata tgtctctttg ttcttacatt agtgagtggg aattgatttg 180
 ctgaattttc attaatgaca ccaatgatgc atctgagttg ggttattgat tgtgatcaaa 240
 ctctggtgaa tactgattga tgtaagcttt tgagttaatt aaggtctaag gtcaatcttc 300
 taatgatttt gagctctcca acaataccaa acaaccgtga aaagtgggtg ggtacctata 360
 aaagatagag gcaaccctat gaggggggag gtgggtgatc ccttttcttt attttctggc 420
 gtttccttgc gcttctgttt ctcttactct cgtgttctcg 460

<210> 359
 <211> 254
 <212> DNA
 <213> Glycine max

<400> 359

tagcataata taaaaccta ggaaacacag attcagtatg ggatacatat atgatatgac 60
 atgacaagaa acagacaata tggcacattt tagaagttat acagatatga tatgtatctt 120
 aattctaaca tggctacatg acatgaccac tggtttcaag tgtatgtact tcttatttaa 180
 gaattatgag aggggaaggtg tttatcacag atatgtggca ctatcagata tacaaaagta 240
 atgagctatc aatc 254

<210> 360
 <211> 114
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 360

gccttgctct tctctactct ctatagtgtg gtcatagatn tatcatntaa cgcataatat 60
atagagaaat ctaactctgg gttctgatat ctgaatcgaa tgctgacatt actc 114

<210> 361
<211> 456
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 361

caagttgtn tgataatgac aaagatgatg acaaaaagcc caaagaatga tttcaagaat 60
gagtcaacaa gttcaagatc aagtttaatt tcaagtttca tgagaagaaa tcaagaagat 120
tcaagaatca agagaagatt gatttcaaga atcaagagaa gattgatttc aagattcaag 180
agaagatgaa ttcaagattc aagagaagaa atcaagaaga cttcacaagg gaagtattga 240
aaagatTTTT ccaaaaacaa catagcacag ttttgTTTT caaaagagtt tttctcanaa 300
natttctaag ttaccagagt ttttactctc tagtaatcga ttaccagttt cctgtaatcg 360
attaccagtg gcaaagtttg gatttcaaag tntttaactg gaattgcaac gttccaattg 420
attncaaaat ggtgtatatc aacacaagat attggn 456

<210> 362
<211> 442
<212> DNA
<213> Glycine max

<400> 362

taattgttta attcttactt cttaaatgta cgttatatac ttgttatagg aaccttataa 60
ttctaagtat atatagttgt agtatgggtc tctgccttaa ttgcataggt agtatgggtg 120
tttgtgattt cttgttcata gtgatgctaa tactctatag ctggatgact catatcaagt 180
tatatttcat aaggaatact cttttgatcg taccttctaa ttctagtgca acctatcttt 240
ttttgtgttg cgtgcttaag tcaaataaat ctgattcact tgaaagcctg agtataatta 300
attctgtgtg ctatgagact acatcacaca atgggactac ttgatgcttc tatcacaatc 360
aagtgattgc tcatgtctta tacgatccac cttgcggtca tgtcttgctc tcgacttcac 420

gcaatctgct gttatctcaa cg

442

<210> 363
<211> 358
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 363

gtatgtgtat gcagaagatg ataggatgaa gctaacttac aataattcca tgtgctcctc 60
tatagtaact gcttgnata gtctgaatc gctcctgtcc agctgtatcc cactaaacaa 120
aacacaacan atgaatccaa gcttggcatg catgtggcaa taatcatgaa tatgatcaag 180
tgaaatgatg atntcgctnt gattctcatt acccaacacc gacagaggtg aatgcagaaa 240
ttatggtaaa ttacaccana caaactanaa acacaataat atagaatcgg ggcattatan 300
atgaagtagt cgggtaaaga cattgattcg tgtcagtga ggtgatact cacaatct 358

<210> 364
<211> 529
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 364

tgcttgaatc atcgaancct ggatcctctg agtcacctgc ggcacgcaag ctctcctcta 60
ttgttgatag agtgtgacat ctatgatcct catagtctcg tgctgtgctc agtgtacggt 120
tagattttgc gcttaagaaa caactgcagc atttgcagct gatattttcta tcttctctct 180
gcatttttct tacttgaagg acattaacaa aagccagttt ttttttgtat ttttcacaca 240
tttttgtttt ggtatttgac atttttgtcg cattaattaa ttatattata ttaacataca 300
attagagcat gagagatgga atctattaac agtttcaaaa ttttagtttg agcctgacag 360
caccaaaaag acagaaattg caaatgaaa acagtgctaa tattttcttt tcttttttgc 420
acgagacata tagttgaatt ggcagtatat ttgcagagaa tatctaaata cagctaaact 480
atctcataat cttatctata atctcgata tagacagtat agataggct 529

<210> 365
<211> 376
<212> DNA

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<400> 365

<210> 366

$\langle 211 \rangle$	435
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<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 366

taactgtctt	tgggcttggc	ggccatgctc	aacaaagtac	tttcgacacc	tactgtacgt	60
tgatttcacc	aatgctgata	tgggaatgct	gctataatct	ctatcaaata	ttatcgatcc	120
atgtagatag	atattattcc	tgggtcaacca	acgctaaaaa	ttactgtcta	attaatcctt	180
gctccttcta	ctattgctag	tttcaccata	cttttccgta	aacttaatcg	atgtatatgt	240
ttcgcgctgt	catagctgan	tcacagccta	agtcacgtat	agacaatcta	agatgcatgg	300
atgatagccc	tccacaagat	aacgatgact	atgcttatgt	ttacctactt	atcttgaatc	360
taatcgaatg	gtgaatgact	actatacatt	ctatagttac	ctcaaattaa	cggcaagctc	420
gtctgcagac	taacc					435

<210> 367

<211> 468

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 367

ttgagaatca atttgttggc tcagtgtgtg attttcgtac aagccattct tttgttcttt 60

atgttaaagc ttgcattctc aacgttntga aaggaagatt ctgtttttat ttctttttct 120
 tttcccttac gatcttatgc ttctcttgac tctgcatgct nttaactntn ntcttttttac 180
 tatatactga tgactgtact ttcaatagta ttattaaagg gtaatttctt ccaattaccc 240
 ttattaatgg ttaccttcca tctcaggatc aggaanttaa ttaataacat gtttcattcc 300
 cacttaatta gcanagntnt ctgaattaac anaaggtana agggactatg tttctttttg 360
 tctctcttta cacataagag agtatcctgg tcaataactca actcacgtaa taaagttctt 420
 attgaataac ctatgacacc ctataatagc tttatttaga tatctaca 468

<210> 368
 <211> 351
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 368

tgteccctctc aagagattnt ngattgattt ctagtttgta agcaccatc ctcttctcct 60
 atgggtccctt gagtttattt tcttctccca ccaagtagac atganatggn gtttcacttc 120
 anattatgat tggtaggtga aaatataatt ganatgagcc tgagtcacac caattcatta 180
 naatgaaagg gaattgctat ttgcactcct cctttataat aatacaatcc ctatttattt 240
 atatttttcc aaaatatccc taanaataca ttccaatgt tcaactccttg caatnntctt 300
 tcgtcanatc cctactgtga gtgcgagcaa agagcaacaa tacaccatca a 351

<210> 369
 <211> 334
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 369

tagaattcaa ggccctgtaa tccacacaca tgcgtcaagt tccatccttc ttnttcacga 60
 gtaacactgc ggaggagaat gggcttcgac tgtgttgcat tacactggaa tngagtaatt 120
 ccataactgt cgttcaattt ccgacttctg gaagtatgga tatttgtang gtctaacata 180
 tngttingcct tcaaaacaaa ttatctgctg attaaccaga aatgagaaac gagcaactca 240
 ttatatatag ggaaataacg ttctgaagat gattcaatta tgaaaaaagg acagattaac 300

cgcatactat tttacctctc tcttcttctt tgtg

334

<210> 370
<211> 215
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 370

ataatcatgc atatgagcat ntcattctca aaagccaaag actaatgagc tggtatcatc 60
attagaagtt aatattgcat gatcataaca agagagatag tgaaatacca gctttaccat 120
atgctgcatc agcaggttgg actttggagc caacagcagc agcaccaatg agcacagtta 180
tagctaaacg gcgagagatt gngctgacat caata 215

<210> 371
<211> 416
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 371

atgacgttat gttttgatcc atttntttta attcatatta aggggtgtaaa ataaagtana 60
aagtgtatga aatttttatc tttnttggtt anaatttctt ttaagttaan actggttcga 120
caattttttt ttataaagtt tagaattatt taaaactaac tgcactactc aacannaaat 180
atattgataa aataaacaac actaagntaa attttttnga aaaattatgt aaaattaact 240
agtaatttag ttgtgcaatg cacaagattg taattaanag gtgaaaaatt gaacaggngt 300
tctaaacaat cgagcactat nttcttaatt ntatcatccn gatacaccac atctaaataa 360
aaggctaaaa atatactata tatactcttag atntctttca tctccatgac atgact 416

<210> 372
<211> 441
<212> DNA
<213> Glycine max

<400> 372

tggagaatag aaagtcgcca caaacattag tatggaaaga taaacatggt atttaggggt 60
tttgtgcaaa tacaaggaaa aatgctatta ccattcttgc cctcttagcc tcttcatcac 120

tgtcattacc atcatcacca acagctttcc tggaaaagtt catagcattc ttaaattgta 180
 agtacatcaa taacaatatg catatcagaa aagtgaacaa aaaaaacaaa ataaagatat 240
 tggacttcaa accttctttt tgcaatggcc cgatcatcat gaccagcctc agcatggcca 300
 tggccatggc tgtagtcagg aaccctgcta acaacgtccc tcagaaagca aagacattat 360
 agcttgacac aatgttttct gtcatttcat tcccattgt acgcttatag ttcgcagatt 420
 atagagcttg aaacagatac t 441

<210> 373
 <211> 333
 <212> DNA
 <213> Glycine max

<400> 373

tattctatat ttgtgatggt tgctttgcag tgtgtggtgc tgaatacatg atagtcgtgt 60
 tattgattac aataattatt agtctggggt taattttctt tatgtgcgct attcacatga 120
 gatgttcgct atgaatgacc acatgtttgtg gaacaatgtt aacaatgtta gttatattgt 180
 tccgctcgag cataagaggg gtgcaactag acctcaaac tactgggaga ggaaccttat 240
 gagaatctca atcccactat attgaggaaa caacaacaag caagtatcaa catagaagat 300
 gtgggagaat taaccacaat gagagaaagt gtg 333

<210> 374
 <211> 592
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 374

tacacgtcca ctaccatata cgatgcgtca atctcanatc tgtctaacgt attaacctac 60
 annncnnacc caagacaggt tgattgatgc antctcgtag ccgngatcct atacagtnga 120
 cctgcatgca cgcaagctta tgcatagcat agagaggcac ttgggtgtact tgcaactcgc 180
 cctagagctc gccagggcaa gctgtttgctt cactctgaag taacttgggt cacataggtg 240
 agctggttac tttagcccta agccatttgg ggggtgcaggt gagttagagg ctagcctgtg 300
 cgagccaggg cctagaaaat tggcttaaata gacccttttg cccctcccc ttgagtagct 360

tccgcatctt tgacaaaac atcgaatgat ctttcgtctt gcgcggtaac tgggtgttgaa 420
 caactcaatt cagctatcga gaatcacata tccatgaatg atagtccctg cacgaactta 480
 ggctgacag tgcccccttt acttatttct atcggataa aacgaagtca tattaggcac 540
 tattctattg agtgcgctgc tatcactggt caccggcaat ccatggatat cn 592

<210> 375
 <211> 347
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 375

gctgctcgcg ttatgcgaga cggagaccaa catgctagct attatcgcca agtaccttta 60
 agagataggt ctagccgcgg cccacgagca taggattgcg ggcgaatatg ctcaagtata 120
 cgcggaataa gaggctagag gaagggatg cgactctcta caccaagagg caaccatgtg 180
 gatggatcag gttgctctta ccttgaacgg gagtcaagaa cttccncgat tgtagccaa 240
 ggccaaggcg atggcagaca cctactccgc ccncgaagag attcatgggc ttctccgcta 300
 ttgtcngcat atgatacact taatggccca catannatag aaatcgt 347

<210> 376
 <211> 461
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 376

tcccatagga agggcttaat tntcttgagc agtttgtaaa acaactttac cttttcgacg 60
 aacttcaaga ggaacctgga cagggatgct agcctaccat ttagtatttg gacttcttgg 120
 tcattggtct ggctgcacat ctccaatatg gcattgcatt cattgggggtt ggcttcaatc 180
 cctcggtgag tgatcatgaa gccgaggaa ttgcctccgc ctgccccgaa agtacatctt 240
 tgaagggtga ggcgcgatgc atactcgcgg agctcctcat agacttcttc taggtctgac 300
 acatgttggg ctatgctttg agacttgaca accatgtcgt ncacatatac cttgacgttc 360
 catttgatct gctatatana gacttgggtc atcagtcgtg ggtatgtagc gccttgcatt 420
 ttaaagccga agggcatgac cctatagcaa anactggcat t 461

<210> 377
 <211> 295
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 377

cttgcgtcct atccttcaac cacttatgat agccaccaac gacanctggt gctgctcccc 60
 taagctcctt atctcttctt tccactgcat tccacgcctt gcggattctc tgaagtattt 120
 ttgcgttgcc ttcattgaaa ccncatgcaa cgaaaggcgc gatgatctct tccaacggtg 180
 cccctctcat agggtaacct agttgtctta tggctagtat gggattataa ttaatacaac 240
 cccttggtcc catcaagggg tcattcgga atccttcaca tgagcacaac acttc 295

<210> 378
 <211> 434
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 378

atcaatatca tttgtattct tgcatacct tgtatcagat ttatttactc agtagtataa 60
 ataccaagtc agttagaata tgttcatacg gaaattgaat ccagataatc atattatcaa 120
 gtccagtcag aagatgaaaa taatgtattc tccaatgggt ctatcatgat agataatttt 180
 aatgtgatgt ttgagtcgtc actatcacct agntgaagga ttgggtccgat acatgtacat 240
 acgttcgtca tcatagagag tatatcgca cgtcatacgc ttataatact acatgacggc 300
 atgatccggt tgactaatgg tctgataact ctcttatct atttattatc agtcacgta 360
 ctagtcggtt tacagtgttt tataatctga gtgtactgca agtaactcta tccataggat 420
 gaggcattga acgc 434

<210> 379
 <211> 461
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 379

ccttgaacgc acataccggt gccaccggag acccccacgt ggtccctcgt gtcttgcacg 60

cataatatat cgagacgctc ganattgaac aacggaagct ctcgacanat tcgaatggca 420
taactttcac a 431

<210> 382
<211> 452
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 382

acaattgatt ttgaatttca acgttcaa at acactggtaa tcaattacac cattttgaaa 60
tcaattggaa cgttgcaaat ttagttgaaa acttttgaaa tcaaactttg cccctggtaa 120
tcgattacag gaaactagta atcgaatacc agagagtaaa aactctggta acttataatt 180
ttttgagaaa aactcttttg aaaaacaaaa ttgtgctatg tttgtttttt gaaaaatctc 240
ttcaatactt cccttggtgaa gtcttcttga tttcttctct cgaatcttga attcatcttc 300
tcttgaatct tgaaatcaaaa cttctcttga ttcttgaaac tttttgattt cttctcatga 360
aacttgaaat taaacttgat cttgaacttg gtgactcaat cttgaaatca ttctttgggc 420
tttntttcat cattntnggt atcatcacia ct 452

<210> 383
<211> 270
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 383

ggagatgcag cggaagacan aggagaagag gtgagaggag gcgcattcac aagggaataa 60
gccatggaag aaggaaacttc ggcaccaaga atgtgccttg gataagaagc ttggagagga 120
tgcttcaatg gagganaaga aagagagaga gaaagagaga ggggggagca tganaatgaa 180
ggaagaaaag agggagagaa gttgaacttt gaagtgtgtc tcacaagact ctcatcctc 240
anagttacca caagtgttac acatacttct 270

<210> 384
<211> 173
<212> DNA
<213> Glycine max

caatagacaa gttatctccg accatttaat tag

453

<210> 387
<211> 339
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 387

cttatgttga gaggttaagt ggaaaacagc aagatataaa caaaccatac actntttcct 60
ctgttctctc cttttaagac tcanaattga gctgtctcct atctnntggt ttgctgaatc 120
agagattctc tccaccaccc ttntcaagtc acagaatgaa gggtagttag ggaaaataac 180
aatganacaa atctaaaaat gaaagaaagg acaagttggt atgtaatgtg aagagatgag 240
agaggaagag ccatatctca tggcatatat ttctgcaaca taaccccact tatgcttcct 300
tatatgggag tgagatcata caacacctca actactctc 339

<210> 388
<211> 458
<212> DNA
<213> Glycine max

<400> 388

ttcgaccttg gtgatctttg actccatgtc atcgaattgc atgtccactt gtaactcaag 60
agcatcaacc ttccaccaac aaagggttga agaccatcaa acctatccaa aaccttttga 120
agaagagagg aatcttctcc accatgtaaa tgtccttctt catcaatggg ttgagcacc 180
tttttcaccc aagagccatc atgctcttta cgataaccaa aggatgcaat catagtggca 240
ccgattaaga aggatctctt gattggaaca taagggtcag aatcaggagg gatgttatag 300
tgtttaagga agagagtga taggtgtgga tatggcaatg tagcatttaa tcgcaatgcc 360
ttatgcatgc gatatcggac taagtgtgcc caatcaatgt gtcggccttt atgaaaagcc 420
cacatgacaa taagatcttc ttcagagacc tgtgcaag 458

<210> 389
<211> 384
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 389

gtcacaccta ctcattctaa aacttaattc cattccanaa cgaccatata tagggaccaa 60
agtacaacat tccaaatcac catctaaaga aaagttcaac ggtgttctac atatgttcca 120
accaagcaca cacagacaaa catgtcatta acacaaatta taagcaaaca aagataggaa 180
gaccgcgagg gggaatgagc gaggganaat gaaccttaca aacgatgaga gagtgaagct 240
attgtgaggg cgagggcatg caatgatgac gacgataaca cacacgagct tcgacaacaa 300
cactggacaa cttcgacata gacgctntnt gtaacatccc attttttcgt anaaataaat 360
atagagcana taaataaata aata 384

<210> 390
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 390

gaagaagaag aagaagagga aaagaggaag aaagcctctt gcatatgcaa aatctgaatt 60
gtgaggatta gggaagggtt tatgaccctt ataattctcc ttttcacacc catgctttca 120
ctcaaaacac ccacccccta acacacatca agaccatca actcccaaac tcattgaaca 180
ttatgaaaaa ccaattaatt aattatgaca tcaccacata aataattatt tacttcaacc 240
acttaattta aatttaatta cacaggataa tttattaaaa ccaattaatc aaacattatg 300
aaaaacacgg tgttacaatt ctccccaaca agaaaatddd catcctcgaa attttcttgt 360
gaagaanata tcgtangcac tgattaagca cacaactatn tcgctttcta ggggtatgga 420
tctttcctct atgatatnct ggatgggan 449

<210> 391
<211> 179
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 391

gtcgaagaac gggcgaaacc ttcgcgaaat tcttcacgga aaacgttacg gaaatgtttc 60
ggaagcgcct cggcttagat tttcttcacg gaaacaatct ttccaagcan attcgaaaga 120

gagagaagtg cctaaggggc tgaacccctt ccttcttcac ttctccctt atttatagc 179

<210> 392
 <211> 308
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 392

atcttcagtt tcggagttgg tcccctagta atntaattca tgaaatgggt cctnctattn 60
 tgtanaatcg tgcaatattg atcaccagg ccacaattgg acgttgaccg ttagcaagtg 120
 atgttgactg tcacgttctt attagatgat gactgtcaaa gtatcgtggg ctgtanagct 180
 atcttatatg ttgaaagcat gatgttctac aaacatgaca ccaaccagtt aaatcacttt 240
 cacatgatan naaagtgata tntttcgtn tcttaattat gcatggataa ntcttatcat 300
 tcactact 308

<210> 393
 <211> 419
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 393

taatgatggg ccgagttatg ttggggaacg gttacgaacc cggactgggt ttaggcactc 60
 actcacgcaa cttaacttgt ntgacatagg ctaaccgtct tgctgggtaa attggatgac 120
 tgtctaagac tacatcttcc tttccttttc gctcaatatg gcaacgacta gttaattgta 180
 tcgtaacgct cggcaacctc ctaccgaaa gttctctcat tgcagtaact actctgtccc 240
 tctcattcca cttaatcttg cgtaagtaac tgctgatct cacttcatta tccggcacct 300
 aactacatc agtgtcggat agcgattcta tagacggggg tcttggtgc ggtattcgta 360
 tgtccccctc ttttgttttc actatcaacc ccactattct ctcttctcta ctgtcttct 419

<210> 394
 <211> 283
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 394

tctctctctc tctctgtact atatatatat atatatatat atatatatat atgtgcgtgt 60
gtgtgtgtgt gtgtgtgtgt gttaataaaa agctaagtgc tgagtgtgat aattntctcc 120
actcatctca aattaagttg gtggtatctc aaatccttaa gcaatgtagt cctanattnt 180
caacaggctt aatatgagag anattcctac aaacagaagt atattgtcaa taattntatt 240
acacataana ttagacagat acatactagt ggtggtccac acg 283

<210> 395

<211> 116

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 395

accgcgatct tagcaccgag ctccagctgtg gacaaacaat agttcttttt tacttaaagt 60
tttccatn atatatggta ggtcagaatg tgctttgtta ttaccttacc atatat 116

<210> 396

<211> 478

<212> DNA

<213> Glycine max

<400> 396

caatggctta gtgaggatgg agaggtgcaa gttaggaagc atgtagagtt ggatatttcc 60
attggaaagt acaatgataa ggtgctttgt gatgttggtc ctatggagcc cagccactta 120
ctcttgggga gaccatggca atttgataag agagctaatac atgatgggtt caccaacaag 180
atctctttca catatcaagg caaaaagata gtgctcagac cattgagtcc acaagaagtg 240
tgtgaggatc aaagaaaaat gagagagaaa attcttcaag agaagagaga aaaataataa 300
gagagccata cacttgagag ttcaaaaagt caggactaaa ttagggaaac acatgacagg 360
aaacggatga ctgtatcgct tgtagtggag gacacttctt acctctacta cgatatttgc 420
ataggatcat gatactactg gtcattctgc tcccagtat tctcattagt taaattcg 478

<210> 397

<211> 436

<212> DNA

<213> Glycine max

<223> unsure at all n locations
<400> 397

gtgctcctta aacctccatt aattntttgc tttaccttct cttccattgt tttttcttca 60
ttttttctcc atgtatctcc tcaaagtct tgtgctaaat tctgttaaca tgcttcttta 120
gatttttcac ctattaaact tgctatagaa gctaaatttt attttctatg gctcaaattt 180
cttgctcttg atcttgaacc atgaattgtg ttgagtttac gttcctttga gttttgtctt 240
gatatttttt gcggctgaaa actaaaccat aaaattctta caataatatt acagtagaag 300
aaaacctcaa aaatctagag tgacttggtc acctattgta gttntgtcat agaagtcag 360
tctagtcag aaacttgta cataagaatt cttatgttgn gctgaatcnt attctctctg 420
ttctttcgct aactcn 436

<210> 398
<211> 460
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 398

ctcctctttc aaaccatgct atgtgctcgc gactagtcnc tctcttcctt tcgcagcttg 60
agttcactat tgctacccca cagagctccg cganatttat tccggccata atcttctctg 120
cgagccctct tggctctctg ttcaagggtc cttgcggtag ttgcattctc ttcccgtaat 180
ccggaacact ccttcogaat gtgtgtagcg gccaaactga acttctcctt ggcaagtttc 240
gcctttccta actcactntt gagagcttgg acttcttcgt cctcttcogg tgcttcaaaa 300
ctctcttcgc tgatgacttn taacttgggt agccaatcta agcctcgtat atgaactntc 360
aaccattcat ggtacccacc aatgatgcca ttacgaatgc ccctaagttc ttgatctttc 420
tttaacnngg gttccatgcc tttatggatc tttgatagtc 460

<210> 399
<211> 446
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 399

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 tccccctttt tgatgatgac aaccacctgt aggttaggag caacaagaaa gaaaaaatat 120
 ctatttgcac atagtttact ccccttgggt ttttcaatgt ttgcttatat gagacaattg 180
 aagatttcat atttttcata tataaaaagt tgtctcataa aaaatagata ttttttctta 240
 ctattttatc ttttatcttt tctctcccc tttgtcaaca tcaaaaacaa atcatgaata 300
 gagaggagaa aaaaaatgtt accacttggt gtaatgtatg agaatcaagt gataccaaaa 360
 ggcattaaac caatcattca atattgatca agcaaaaaca agtatagtaa cacatcaatc 420
 anaaacacaa tcaaaagcaa tcaact 446

<210> 400
 <211> 469
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 400

cattgacaac atgattgacc ccttggcagc cgaagttaa tgtgccaagc tatgcctttt 60
 ggggttattg ttgcttttga aattttaacc aaaaatgact aaagtaggtt taaacaaaa 120
 atggctaaaa tggctaaagt aggtttaaac caaaaatggc taaagtaggt ttaaaccaaa 180
 aatggaaaat tttgcttttg ttaaaactgg taaaccctat cataatccc tagatggatg 240
 tgctaaccctt ccttggatgt gtaatcagag tgaaccttgc acaaagtcca ctctcaciaa 300
 gttaaattac atagtcactc aatgcacaat gcaattcttt gatagataga aattcagctt 360
 agacaatttt catatctcta tatcaaacia aacacatata ccttgttata tatctatgta 420
 tagcacgaaa ttcaaggctt agaattatac aatntataaa ttaaattct 469

<210> 401
 <211> 451
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 401

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 caaaatacat gaaaatataa aaaaaagtcc ctaatacaaa gactactcaa aatgccctga 120

aatacaaggc taaaacccta tactactaga atgacccaaa tacaaggcct aaaagaagga 180
 aaaatctatt ctaatatatta caaagaagag aggatccaac cttgggtccat ggggtcagaa 240
 atctaccctg ggattcatga gaaccncaag gccttcttta gcagctctag cccaatcctc 300
 ttagagtctt ctatccaata ccccttggtg gtaggatng cttcattccc ttcaacttgg 360
 aaaggatntg acctgaaaat ccgaagggtc tcataatttg ggctccctcc ctcgacacct 420
 cgaaaaaaga ataaaacata tgtattagt g 451

<210> 402
 <211> 380
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 402

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 taacactctg gagtanacct tgatcacatt cctctaaaat gtgacagcac aagtgcgac 120
 aatctaacan aaaaaatctt gtcatgcatt ctagaactaa acacatagaa ataaggcatc 180
 attntctcaa agatcatatg ttaaaagttg attgggtgat tgagttcata gatagtgcg 240
 atcaactaac agacgttntc actanaccac ttgctagaga tagattctnt ttcgtagaa 300
 atgaactaga catattagat gcacttagta tagaatgaca ttctatttgc atagtgtgtg 360
 atgcacattc ttactcatat 380

<210> 403
 <211> 114
 <212> DNA
 <213> Glycine max
 <400> 403

ctaaaaactt agttagataa aggttataga tttaggcgac agtggcagcg gcaattatac 60
 aacgcctgcc tacataaaac tactgattct acatagatat ccaggacgat cacg 114

<210> 404
 <211> 393
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations

<400> 404

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acaagtcata cctaattcng attggtctga ctaaacaatg tanaagtatt tatactagtg 120
atztatgaaa aataaattca ttntanatnt gtgtttaatt ntaatttatt gctggtgtaa 180
ataattttac cccatcaact aattaanaat cttaatgtaa ttataagaat ttaattat 240
agcaccacaca atataatata taggaaccag tacaaaaaat tctttttaac attagttg 300
tttaacattn tatatatgtt ttgtaatgta tgctaataag ttgatataga aacactaca 360
gaaaaacact taaacatggt tgatatattt ttt 393

<210> 405

<211> 241

<212> DNA

<213> Glycine max

<400> 405

ttatcctgct ttgatgaata tgaagcctcg ggaagatgga gagataagaa agaggagaa 60
tcatgttggt actgccgct acatggccaa attccacaac taacaatgca acacttagct 120
agatagtcatt ttcataccca ccacctacta gtaagacact tatcatcaca aggccacctt 180
aatcagcaca aagtcacctg cgcacatcta tataaacacc ctcttacact accaaacact 240
a 241

<210> 406

<211> 246

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 406

atcatgataa caaccaatat gaattccact canaggagtt gggcatgtaa aagccaaaac 60
ttcttcaagc tntagcctta agttgttcac catgttgctc ccctatctct aacaacccat 120
gcatgtagtc caagttcaaa ggattatagt atgttgatag tgggcgcata aaccatatga 180
taagggactc aagtctgtta aactctttag acaaggctgt tagaaccaaa gtcaagaatg 240
gaaatg 246

<210> 407
 <211> 454
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 407

gatatccact ccacaaggtt tgaagtagag gagagcttca accctataac gcaacgtggc 60
 ggacaaaagt gggcagtaaa cttgaatgat cgtcattgtc aatgcggaag gtattctgcg 120
 cttcactatc catgttcaca tattattgca gcttgtgggt acgtgagcat gaactactac 180
 caatatatag atgttggtta taaaaacgag cacatcttan aagcttactc cgcacaatgg 240
 tggcctcttg ggaatgaagc ggctattcct ccttctgatg acgcatggac acttatccct 300
 gacccaacca caattcgtgc gaaaggctcg ccaaaatcaa caaggataag aaatgagatg 360
 gattgngtcg aaccatctga gcaccgaaca naatgcagta gatgtggagc cgaagggcat 420
 aacaggcgct gctgtccaat gcaatctgag cgtg 454

<210> 408
 <211> 304
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 408

tcagctatat gnggttactt atatttgtca tggtggatan natttagctt tatatattgc 60
 taattcagct tatagggtgag aatgaagcta cntagctcc tatatatnna accatngatn 120
 ntaatagcaa tatggcactt ntgagcaaga attactctct ataagttntc atatcaagag 180
 tcanatgcta ttgaaatgga taaatgcaca atataattgg tgtgtatcaa ccctaacaca 240
 acaacactac cacaaaaaca cacacnctat gatccacaat tngaaacgaa agggaaaagt 300
 catg 304

<210> 409
 <211> 457
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 409

aaaggatcga agtgggtctg aaaagaggca aatttaataca tcttgcttgg gcgaatgaga 60
 aaactggggc aattgaagaa ggtgaggatg agggagaaac ccatgctgtg actgccattc 120
 ctatacgacc aagtttccca ccaaaccaac aatgtcatta ctcagccaat gacaaaccct 180
 ctctttaccc accaccagc tatccacaaa ggccatccct aaatcaacca caaagcctgt 240
 ctaccacact tccaataacg aataacactt ttagcacaga ccanaacacc aaccaagaaa 300
 tgaattntgc agcgaanaag cctgtaggtt cacccecanat tccggtgtca tatgctaaac 360
 ttgctcccat atctacttga tactgcaatg gtagccataa cccctactan gtttcctcaa 420
 cctccatttt tccgagggtg cgactcgaac acaatgg 457

<210> 410
 <211> 254
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 410

tagatagatg acatcnagcg ataactgagc catgtgaatg ctcatncgtg tcaggatggc 60
 gtacactagc taacacttcg gcagggggag gtcggaatta tgatcgctgg gctggatgtt 120
 gctgagcagc anaatcatcc agatctgagt caggggtggc atgggtgggtgc gcatgatccg 180
 cacnctctt ccagcagcag tccaggcana atcctgcccc ggtatacata gcaactgggc 240
 gatggcctct catc 254

<210> 411
 <211> 455
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 411

aacctatatt taaaataact taatgccatt aacctagga attaaaacaa acttaatggc 60
 tgagtgtaac tgatattgtg gcaaccaaaa gtcaccccca acagccaaca agtcagccac 120
 catttggctt cccaaaaggc tgatgcctat gttgccaatt gggcccttat tacaacttga 180
 actaaagccc ttttagttga ttaacccaaa acatattttt ggtagccaa ctttacaagg 240
 attggggcat tatttagaca aactaaacac tctaaaactg aaataaagtg gtgtcattta 300

<211> 278
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 414

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 gttgtaaatt cnagttgaaa aacttttcan aacaatattg ctactggtaa tgtcatatccc 120
 taatttcgtc cggggacctt tgcttgatga catgcgacct ttctttggtc cttgtgaggt 180
 gcttggcatg catcattang caatntgtga gattccagga catgccgaca aaccaacaaa 240
 atattgatgc acaaatccgt aagtttccgt gacacacc 278

<210> 415
 <211> 457
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 415

cgggccattc caagtgttg agaagatcaa cgacaatgcc tacaagattg acttgcctag 60
 tgagtataat gtaagtgcc ctttcaatgt gtctgatcta tctctttttg atgcagatgg 120
 aggagccttg gatttgagga caaatccttt tcaagaagga gggagtgatg aggacataac 180
 caagggcaag gaccatgaag cacttgaagg tcccatgacc agaggcagac ttaaacaagc 240
 ccaacacgtc atagagacaa ggctgggtcat ttgtatagct gccattgatg atgattgaag 300
 gcccaagtgg agaaagatga agggccagag gcagaggcac taccaagact actaattggt 360
 gttgaaggcc cataactaact tgaaggccca agttaaataa gttnttagtt ataatttatt 420
 tntattggaa ttctggccca tactgtntag aacgcn 457

<210> 416
 <211> 511
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 416

ngtgtgctcc tgctcttacg gacctatgaa tctcagctnt cacaagagtc ttcacaaaata 60
 actatcatga agcaganaac taacaaaact acccatcata tctcccaaaa ccccatatccc 120

acaaatattgg agcttcaatg gagaatgaag aagaagagaa tggcaacgtg agggagagag 180
agagctgtct gaaataatgt ggggctgagt gaagagagag agagttgctt tttgatttta 240
aaaaggcttt ttcctcattt cttattattt tattataaac tatgccacat gtctccattt 300
gagtggagca aaaagggccc actttccctt ttgactgtga cccatactca gccacaaaag 360
tgaggaaaat ctgacctttg aaatgctaaa atcctgcctt ggttggcgtg ccgtttctct 420
ggttccagtt cctcgcgtt ctctgcgtcc atcggngcca gttttcgaaa gtacgcaata 480
tatatatcan aacgctcaga ataaaacccc g 511

<210> 417
<211> 445
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 417

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gatgctgctc tagatcaaag tggcaaattg gatcaaggag aattggtgag aactaaagct 120
aaactacgga ttgcacaagg aaaattaaag aatgcagtgg agacatatac ttttcttctt 180
gctgttcttc aggttcaaaa taaaagtta cgcacagcaa gtaagggtgt gaaggatatgt 240
gtcaatgata aatgtagaat atatcactta gattctttaa taactcanat tatcaacggg 300
aaaggaatat tagaagcttc gctcacagtc ttanactcta ccttttaaca tatgtagaat 360
aatatattac ttttctgatt tatctttcat acagaataan ggaaaccgtg acagaagact 420
ggaaatggaa atatggcttg attat 445

<210> 418
<211> 480
<212> DNA
<213> Glycine max
<400> 418

ctcttatttc ttggtaaagc tctatctaaa taaagttggtt attactgcaa gaaatcagat 60
ttatctccca ccctgtctt tctcaatcat tccaatcccc accccccaaa gtccatgagg 120
gtacccttca ttatgtggtg cttacggctt atacaaatct aataatttgt ggattcaggt 180

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<223>      unsure at all n locations
<400>      419
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<210>	420
<211>	269
<212>	DNA
<213>	Glycine max

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atgttgcana	tatttacaat	agacctnctc	aacctcaaca	gcaaaatcaa	ccacagcaga	120
acaattatga	cctctctagc	aacagataca	accctagatg	gaggaatcac	cctaattctca	180
gatggtccag	ccctcagcaa	caacaacagg	gggtacggaa	agtaccccct	tgaattgtat	240
attcaagaca	tttgagaata	aacaaacac				269

<210> 421
 <211> 427
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 421

ctaanaagag gtatatgtga tgtgagagaa aaaaacatac tcatactccg attattaaca 60
 gatgctttct ggtgttgcac acgagttgcg gcaggatgag ccaagctttc aatggaccca 120
 atcaaaataa gaacattgaa gcttctaaag agtgacacgt atattataca aatataatag 180
 ttagaaatag atagtatcat attatagctg atatatatca gatgactaac ataagatgat 240
 cactgctagc tggacggcag caganaattc atgccaggaa acgattaaat tttgacttta 300
 ttaattcttc tagcacctta taatggaaaa aagagttgat agatttgcgc ctaaactttt 360
 tatttaaaac aaacagagtt tccaacatcg attgagagtt tttttatatac aaacctgtga 420
 ataattgt 427

<210> 422
 <211> 365
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 422

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 actttggaaa aatgccagtt cttggtgtct caactgctga ngctgcacgt gagggatatga 120
 aaacacatga cctcgttntc tccaacagac cacatcgtaa gatgtttgat atcctcttgt 180
 atggttccaa agatgtggca tcttctccat atggcaacta ttggaggcag ataaggagta 240
 tatgtgtctt gcattctctc agtgccaaan aggttcaatc ttttggtgca gtgagagaag 300
 aagaaatctc cataatgatg gagaagatan ngcagtgtgt cttcttgatg ctgtgaatta 360
 tctga 365

<210> 423
 <211> 448
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 423

ttcanagttg ttttgatgat aacaatgatg acaacaaaag atgatgacaa aggtgatgaa 60
aaaaagctca aagatcaaag aacaactcaa gtaaatcaaa gaacatctca agtgaatcaa 120
gaacaagtca agagttctag aatcaagaag aattcaagac ttaagaagaa agcctagaat 180
caagaatcaa gattcaagat tcaagaatca agactcatga ttcaagaatg aagaaaagac 240
tcaatcaaga taagtattaa aaagtttttt ttttaaactt tgaatagcac atgagttntt 300
gacaaaacct ttaccaaaga gttnttactc tctggtaatc gattaccagt agcaaaataa 360
gtttgaataa gttttcagac tgaatttaca acgttccaat tattntcaaa aggctgtaat 420
cgattacaat gttntggtaa tcgantag 448

<210> 424
<211> 469
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 424

ccatcaaccc catgccaaaa tacatganaa tatataaaaa aatccctact acaaagacta 60
ctcaaaatgc cctgaaatac aaggctaaaa ccctatacta ctagaatggc caaaatgcaa 120
ggccaaaaag aaggaaaaaa cctattctaa tatttataaa gaataatgga tccaaccttg 180
atccatgggc tcaaaaatct accctaaggt tcatgagaac cctagggcct tctttagtag 240
ctctagccca agcctcttgg agtcttctat ccaataccct tgggggtagg attgcatcac 300
accatacaac attggttttg accatcaatc actatccctt tgtggttgat tcaccttcaa 360
atcatattta tgtttggaag agagaaattg ttgttggtgt gagcgtaact tctcattctt 420
tgttgatctt tcacactcca ccttcacctt cactaatcaa ctcaaacct 469

<210> 425
<211> 234
<212> DNA
<213> Glycine max

<400> 425

tctgtgacac catcagacct atgccttcat gcagaacctg agcaataggc agccgaagtt 60

atgctgaaat atttacacag acctctcacc ttagggcaaa tcaccatgca aacaatatga 120
 ccttcagcac agaacaccct gatggggaat acctacctca aggtcagcct cacacaacag 180
 agctgctctt cttcaaatgt gtggccagag acatcattct cacatccaca caca 234

<210> 426
 <211> 418
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 426

gctacaacaa ctgcagcaag acccactatc cacaatgaca gaacaatatt tgtntatacc 60
 ttgcatcttg tatgaaagat gttctctctt tgtgtttggg ttaggtcaca agattgactc 120
 ccaagtaacc ttctgaccat tagaagatca cttcttcat aggggtaaat ctcttcaata 180
 tgggtcatcac cattggcttc accctcactt ccactngagg aaggagaaga tgtagcctnc 240
 ttttggctac tatagatgtc ttgaccgtc atgatcatgg ttttctttgt ggggcaatga 300
 gaagcaatgt ggcctagcaa tgcttgttct ttcctcctcc ctaagtctag ctctcagaag 360
 ggagtagtgc atttgtggct atattcttca cactcatact cccttngcta agcttttt 418

<210> 427
 <211> 589
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 427

ctaacgggctt cctccatag tcataggcgt taaagcgggt gtaatatctc gtaacacttc 60
 aaaanncccta gcactagttt gaatcatgcy tcgctgagac centagagtc gacctgcagc 120
 atgcaagctc tgtccttaga tcctcttctt tggactatac tcataccaag caacattatt 180
 gtacaacata ttataaccaa cacttaatcg gcagattcct cttagcagac taagattcaa 240
 ttctgcttca ttcaaggctt aaggcaacaa tacattttcc aatgcttaaa tcacctaacc 300
 gggcacacaa atgggttgatc agaccatgag catacaaaat ttaagcactg aaagaagcat 360
 tgaacacact agaaactcaa tcaattagat attaaaataa ttacatcagg tgttcttttag 420
 aaatacccaa caagggtggt tagcccacca ttacagacaa acccctatca ataagagat 480

aaataaacgg taagatttct tgaaagctgt ctttttgctt ctacagagct ttttccaaaa 540
ggcacttggg tgcttataat ttgtgcaaag cgtgttaaatt ctggagtan 589

<210> 428
<211> 192
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 428

caaccaaaga tatgaagatg tgagatgtnt ggttctctgc catcgaataa ttcatatgaa 60
gtnttctnta aaatgggtct tattaaagcc ctatctaaaa tgtagcatgc agtggttaacg 120
gcttcagccc ataagtattc tggaagagga gtatcattca ataaagntct agcaatctcg 180
tccaaagatc ta 192

<210> 429
<211> 526
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 429

cgttgatcgc tcgtgacgag ccnnttgagt aacatcgacg catgcaagct tgcatatgaa 60
gaanatgggt gggtgggcat taaatgtgtg cattaatgc atatactttc tcatgctaag 120
aaactagtct ttgtcgcagc gtattgaaca cttaaggagg aaaccacttc ttttgtgtta 180
gaacaagttt atgcaacaaa gttcttcttt tgatggcgat tgagaaaattt tagagcttga 240
cttcatttat tctcatagg atgtgacaaa tcttaggaga atatctctgt aaaatagatc 300
tcaaacacaa gagtattaaa tgaagtctta catgtcaact ttaatgttgt atcagatcat 360
gatttcattc tggctaccat tgaaacatca gatgcagact ntgcaaaaaca tgatttgata 420
gcacataaga tgcaactnta aaccttcgta tttgttttca tctaacaaca tgtcttaaga 480
cataaatgtc ttttaaccta gcaggaatgg tattcctatg aatacg 526

<210> 430
<211> 250
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 430

acttcatata gcagtctata tggctgggta ttngaaata tcacaattca tatgtcaggg 60
aatcttgtca cctagagaaa attgagaaag attntcttag ataatagtcg agaatagaga 120
gtagagacgt tgggaacaga gataatatgc tctgtcttac tgcgacaaa actacaaagg 180
gtgtacaaag gagcactcgt ctttcacacc acgatgagtg ctcgtcaca cactgtacac 240
atgttattac 250

<210> 431
<211> 459
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 431

acatgtggta ctaggtggcg gtcgggcat gatgcacaac aagctttcca catccacaat 60
gcgcgcataa acccaccatc cctttttgcc cacctccaac tgagctcag tactcccacg 120
tagcccatat ctcgtttct ctcaacaccg ggtaccatc aatcctcca agcttcaca 180
acatccaagc aaaacaacat tcaaacagca caagctatca cagccaagca aaacagagca 240
aaggcagaaa actctgctca acacatcaac caaatcaca gcttttctca cttaaagacc 300
acagtaacaa ttccttcgat ccaattcgtt aaccggtgga tcgactcaa aattntactg 360
gaagtctata gtgcataagc ctacattntg accgttggga tctactggca nacatccaga 420
actcattctg cactactctn tccacaacca gcanaaach 459

<210> 432
<211> 201
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 432

agctacctag tctataaata gaagcatgtg taacactagt tgtaactntg atgaatgaga 60
gtcttgtgag acatacttca nagttccact tctctccctc ttttattcct tcaatttcgt 120
gcgccccct ctntctttct cttctctnt cttttcctcc attgaagcat cctctccaag 180
cttcttatcc aaggctcacc t 201

<210> 433
 <211> 456
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 433

gaaacttcct gcttttattc gttgaccaca gagtgggtacc tggagatatg tcgcgngngt 60
 caggagacct tngggacgtc aggtgggggtg ctattgccca aaaccaagct tgaccaatcc 120
 cgaccaacc cgggcatagt cagtcagtga gagcctgtga tgtacctaaa caggcgagct 180
 cctggcagtc aacagataaa aggaacaaaag accacaaaagc aaggaggctt gtgggtggctg 240
 gccagctgtg aactttgatt gatatgtgag atttggcctc tggtaatcga ttaccaaggg 300
 tgggtaatca attacaaggc ttanaaatga agacagaagg ctaagatggc ctctagtaat 360
 cgattaccaa ggggggtgtaa tcgattacca ggcttgaaaa cgaggtcagg aagccatgan 420
 ggcttctggt aatcgattac caaggggggtg taatcg 456

<210> 434
 <211> 318
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 434

gaataaagag ggggagaagt agaactttga agtgtgtctc ataagactct cattcatcan 60
 agctacaaca agtgttacac atgcttctat ntatagacta ngtagcttcc ttgagaagct 120
 ttcttgagac aacttccttg agaagcttct ttgagaaaac ttccttgaga agctagagct 180
 tagctacaca caccctctc ataactaagc tcaccttctt gagaagcttn cttaagaaga 240
 ttcgtanaga agctagagct tagctacaca tacctctcta atagctaagc tcacctcctt 300
 gagatgagaa gctagagc 318

<210> 435
 <211> 215
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 435

tatgtggact aggtggcgat cggacgatgg tgcaagtcga ctcttcacat ccacaaatca 60
cacataaatc catcatcccc agntggccac cttcaactga gctcacgtac tcccacgtag 120
ccncttatcc tcgntccttc aacaccgggt gtccatcaat ccctgcaagc ttccacaaca 180
tncaagcaat tcaacattca tacatcatga actat 215

<210> 436

<211> 308

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 436

gactganaat gttattcagt ntgtcagaat ggatgtgaag cttttgattt gttggcagcc 60
caacctcatt acanacctag aaagtccttc ggattcattn tgtgtgttta tttctgtatg 120
gtatgagatg atatgcanaa gttgggactt ctgttagttg tttataatgg aatgatccta 180
aacacttgag cttgagtga acaacgactg tgaggctntg gttgatgatt ctttccttga 240
tatctgccat tctcactaac ttattntagt tgtgactcta atgcatatgt tcctatcatt 300
gaaaagct 308

<210> 437

<211> 456

<212> DNA

<213> Glycine max

<400> 437

actaacgtcg tcttctgcga cctttgtcaa tcgcggccga caagcccgtt gacacgtgga 60
gatttacgtc atcttccgag ctcacaagat ctgtcatact gacttttgag tcacgtgac 120
ggccggaaat atccgagtgg ttatccgtat aaactttttg ctgtctgtaa gacgaaaagc 180
ttgatagcac gcagagacta acgtcgtctt ctgtgccatt catcaatcgc ggccgacaag 240
cccgttgaca cgtggagatt tacgttatct tccgcgctca caagatctgt catactgact 300
tttgagtcac gctgacggcc ggaaataccc gagtggttat ccatataaac tttttgctgt 360
ctgtaatacg aaaagcctga tagcacgcag agactaacgt cgtcttctgc gaccttcgtc 420
aatcgcggcc gacaagcccg ttgacacgtg gagatt 456

<210> 438
 <211> 254
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 438

actcanaagt cagtatgaca gatcttgtga gcgcggaaga tgacgtanat ctccacgtgt 60
 caacgggctt gtcggccgcg attgacgaat gtcgcagaag atgacgttag tctctgcgtg 120
 ttatcaagct cttcgtctta cagaatgcan aaagtttata cggataacca cttcgggtatt 180
 tccgcccgtc agcgtgactc anaagtcagt atgacagatc ttttgagcac cgaagatgac 240
 gtanatcacc gcgt 254

<210> 439
 <211> 487
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 439

atgtataata aaataggtaa tgtgatacat agaaggatga caagcaataa aatcgaacat 60
 aagaaagtat tacaaaaagt acgaataata ttaagataac ctacataat ggagaaacag 120
 cattcttgat aacaattcac ttccatcaca agaaanaaga tctgataccg tggactgac 180
 aaacgcatat tnganaaaga tatagaatag ttatatcttt gattcagtgt atggccaaaa 240
 attgacggta cagaatgtat gaagagagtt tagtctaatt aactaaacag aatataccaa 300
 tattgtaaac tntagtatgg tggtcagcta gtacggataa ngaaacaata caaaatttga 360
 tctaaataat atagctctta tgtcaaagca caatangatg atttttaaca aatgactgaa 420
 tcaacacgca tatatttatc aatctccaca aagatagaga tcatataaac atctcttata 480
 tttatat 487

<210> 440
 <211> 439
 <212> DNA
 <213> Glycine max

<400> 440

ttgtcgtgat taccaagtga cagaacaaca ttacctgtat tgagcaactt caataagaac 60
 ctttacaggg atgctatcct accctttagg atttggactt cttggccata ggtatcgctg 120
 cacatctcca atatggcatt gcatttattg tggttggctt caatccctcg gtgagtgacc 180
 tgaagcccg gcacgtgcct tcgtctgcac ctaaagtcac tgttgacagt ggcagcccat 240
 gtatactccc ggacatctca tagacctctt ttagcgccgc aacatgtagt cctctgcttt 300
 gtaattcgac tcctatctag tgcaaata ccttgacgcc acattcgatc tgatatttat 360
 aggcataaggc tcaacctacc gtgtgtatgt caccgctgac ttatctagcc ccacgcgcta 420
 ccctaaacaa cctcgcgcg 439

<210> 441
 <211> 419
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 441

caacaagtat tctncatcat gcacacacac acantcatat caagagaaaa aggacattaa 60
 tccttttccc ttcctanggt cacaatcaac atggcccttc aagtggaata atccacttta 120
 ccaaatacaca caccaccaca tatccaatca ccaaatacatt actagacatt caaagtanna 180
 tttttctgaa ggttgacac nctttgacct aacctanag tgcgacgaat ctttaattaat 240
 atcattaata aactcatata cataacacac aacattcttt accaagtggg cacactcaat 300
 tgggtcttaan acatatacaa tagcaattct tataatttca tattataaag tctncatcan 360
 agtanacaat acattctaca attcacatca ttaatttcat gcatttcata tacattcat 419

<210> 442
 <211> 463
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 442

tcaaagcctt gtatggattg aaataagcta taagagcttg gtatgaaaga ctaagtttat 60
 tcttactcta gatagttact ctagagaaat agtggacact tcactattca gaaaggctta 120
 gaaaaaggat ctgctgatta tacatatata tgtgaatgac atcatTTTTT atgtaacctc 180

<212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 445

 ctccagctaataaaggaggaagtgaggaggttggaagctactttggccaccgcccagagggga 60
 gaggaatgctgaaaaggaggagaggattgnggttgagcaa aaaacatataacaatgttta 120
 tgaagcttatgagtgtgggttgaatcactgcatctaacaattacactttgactatgaggt 180
 cccaaatgaa tccattttcaacatgaacaaagatgtctataatggagacttaatcttgat 240
 tgatgacatttcggatgagg tgggatcaaa tgggtgggcagccaactacccctcttggtga 300
 aacccaaatgaccaattcggctgaggagga tgtggacgatgtcccatatacgagccanaa 360
 tggcgtctctt 371

<210> 446
 <211> 436
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 446

 tgttcgcacatcgttcgcgtgatatgataccactcgacaa ggtttgaagt agaggagacc 60
 ttcaatccta taacgcaacg tggcggacaa aaatgggcag ttaacttgaa taaccattat 120
 tgtcaatgcg gaaggatttc tgcgcttcac tatccatgtt cacacattat tgcagcttgt 180
 ggttacgtga gcatgaacta ctaccaatat atagatgttg ttacaccaa tgagcacatc 240
 ttaaaagcat actccgcaca gtggtggcct cttgggaatg aagcggcaat tctccttct 300
 gatgaggcat ggacactaat ccctgaccca actacaattc gtgcgaaagg tcggccaaaa 360
 tcaacaagga taaggaatga gatggatngg gttgaaccat ctgaccaccg acaaaaatgt 420
 agtagatgtg gagctg 436

<210> 447
 <211> 442
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 447

agcgcggggtc tgggagacaa aggtcaagcg ttcgcgatat gcgaggatga tattccgagt 60
 actttggatt tggtagcacc atgccctcct gatttccagc tgggaaattg gcgagtggag 120
 gaacgccccg gcatttacgc aacgagcata atgtaaacct ttacggttttt aaaagctcta 180
 tagttgggcc taggcttttag agtttttctt tttgttaagg ctttgtgtct tttgtttttg 240
 aatttataat acaaggatct ttcttcatct gttcctggtc tctacccatt ctcatcatt 300
 tgcattgttta cttctttntc tgaaacggca gatccgatga cgagtcccc gaaggtacta 360
 atacctggga cccgcctatc gacttcgagc gagaaatgaa tcanacggaa gatgaaggaa 420
 atgaggatgt gggacttccc cc 442

<210> 448
 <211> 410
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 448

ngagttgagg aagtgtagaa ggggtgaaact tcttgctntt attcgttgac cacagagtgg 60
 tacctggaga tatgtcgcg nggtcaggag accttgngga cgtcagggtg ggtgctattg 120
 cccaaaacca agcttgacca atccccgacc aaccgggga tagtcggtca gtgagaacct 180
 gtgatgtacc taaacaggcg agctcctggc agtcaacaga taaaaggaa aaagaccaca 240
 aagcaaggag gcttgtggtg gctggccagt tgtgaacttt cattgatatg tgggttatgg 300
 cctctggtaa tcgattacca aggggtgggta atcgattaca aggcttataa atgaagacaa 360
 gaggctaaga tgggtctctgg taatcgatta ccacgggggtg taatcgatta 410

<210> 449
 <211> 455
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 449

tagaagaaca aaattgcctc aatcatttcc aaatatgcat gtgaattang aagcatcaac 60
 aagaatcaag ccaaggctat tgtgcaagca atcaatgggg caaaacacac caaatgatta 120
 tgatgatgga tgggtcaaata tctcaciaag gtaaaactcat cactttcaaa ttgagctttc 180

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<400> 450

<210>	451
<211>	384
<212>	DNA
<213>	Glycine max

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<223>      unsure at all n locations
<400>      451
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<210>	452
<211>	401
<212>	DNA
<213>	Glycine max

<223> unsure at all n locations
<400> 452

tggatttcct tntagtaggg aatctatcct tcctaagatg gagccaaacc cagtcaccct 60
cattaagaac tagctctttt cttcctctat tgcccttagt tgaatacacc tttgtttggt 120
tctctatttt gttcttaacc ctctcatgca acttctttac aaattttgac ctagattecc 180
cttctttatg tataaaagaa gtgtccagtg ggaggggaat gaggtctaac agtgtttagg 240
gattgaaccc atagacaacc tcaaaagggg actgcttggt ggttctatga acccccctgt 300
tgtaggcaaa ttctacatga ggaagatact catcccaaga cttatgggtg cctttcagaa 360
gagcccttat aatggtggat taaaacctat tcactacctc t 401

<210> 453
<211> 444
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 453

tcgtgcatgc cctttcccat ttcttgtatc agtgacttga gttcgtcggt tgaacggccc 60
aacgcattgt tggtccggtt catctcttcg attttcggag ctattgcttt caggatattt 120
tgagggcaga ccaagttttg tttaaagagt gtggctctgc ttattgttcc tgaacaacc 180
aattctacga gcatagcctc tgccatatta attcagagc tagctagtct caataacgag 240
aagatgtata tatatagcaa ccttagactc tgcacaatat aaactcatgc atgcaaccaa 300
gatgcttctt gtatactatt gttaaaggga agtttctttg acaattcatt cccaatttca 360
tattgcttct tatantattt gctgacttgt actgggtagc gttaatatca acgggtgcaa 420
aacgtctttc ttttcttttg acct 444

<210> 454
<211> 453
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 454

tctttcctaa atcaaagtat aaatgattnt atattataag tataaaccta ttaaaaaatt 60

ccgttcaaaa taataagata attttacctg aaattttgaa ccaactccact tgtttttaat 120
gagccaattc caatcttctg gagatacgta ttttggctca tcattttctt taagataagc 180
ttgatgtaaa tgataaagct tattcttgta ttgagattga attttcttca ttgcatattc 240
tctttcctcg ccagataaac taaaagcatc cttatatcaa acgttttcac cagcaaaagt 300
aaaaattaaa tactatcaag atacattatc aaaccaattt gaagatataa ctaatattca 360
aatatataaa taaattagat ataattaacc taattatata ctaacattca tatgtcgcca 420
catttcatct acaccatcct tactaatttc act 453

<210> 455
<211> 442
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 455

tgttcttagt aagatcgatc agacgatgct tttgttcctt ggagagcaac ccgtcttggc 60
cccagaaagt gttgccaatg ctctagtgga agtgatgaac gatcctttct cacacttaat 120
cgcctgttga taaaattctg caccaaatat ccaaacacca ttaaataaac taaaattcaa 180
ataattaatt gtatggatgg attagtcaaa taaaaatttc tattagccaa gatattatag 240
gaatttgta agtctatcac gtgagcagtg ggcttagtgt tcctttcatt aggagattca 300
tgaaaattnt gtcggccaca naattgtgag acttcttaat caattaaatt aaaaaaacc 360
tagtaaaata aaagttgacc agcatanact aatagtggga atataattag tatttgaatg 420
ccaaaatcaa ccacatattt tt 442

<210> 456
<211> 431
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 456

cgagaaagcc cttctgattc tgtttataca tttctgactc tatggcatga gatgcaatgc 60
acagattgga cctcctgtta gttgttatca aagaatagct tatacacttg tgcttgagtg 120
aaacagttgc tgtgagactg tggtttgagc tactttcctt gatacctgtc ttatgattaa 180

cttcatctaa ctgtatagtt cacatTTTgt tctcctcttt gtctagctgc atattctggg 240
 aaaacaagtg ataggtacac attgcttcat cttttacatc atgcaatcaa taaattntaa 300
 tgcatacacc tttgaacata aacactgcat gttntaccac ttgaggacaa gtgagttggt 360
 ctcttttgct tgaggacaag caaaactatt aaatttgggg agttttagt cgatgaatac 420
 gactaacttt t 431

<210> 457
 <211> 357
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 457

cttctcggat atctgttatt gataaacaag atcaagatga gtatagtttg aggtttatac 60
 acagaatctc caagattggt tatagtacaa gcggttagta aacacagcat atatgatagc 120
 tggcaagcac atggtgagag gtgaatcggg aatcagtaat cacacagtta agtgaaatac 180
 taaagtttat ggagatagtc cgaaatctta cagaaagtca aaagttctta taatagtagt 240
 taaataaata aataaataaa agtagaattt gactaagaat gaaaaagtaa tattttgttc 300
 aataactcat ggnggtatat atacgagtgt tctgacgcgg gaatcacaaa cagaacc 357

<210> 458
 <211> 479
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 458

ctattaggac tcttgcaact cagctcgtac atatagtttc aatccgaggt ctttcaagag 60
 acctagtaaa agtatcagcc tattgttcat tggaaccaac aaagtcagtc atgatttccc 120
 ctgacaacac ctttactctt gcaaagtgac aatctaTcta tgtgtgttta gtttgttcat 180
 ggaagaccat attagatgca atgtgaagag cggtttgatt gtcacaaata agcttagtgt 240
 cctgagtgcc taagtcttgt aatttcgcat gtagctgcta ccatggcaca atgttaagct 300
 ttgatgctga atctagcaac tatattttgc ttcttgtttc tccatgagat caaattccct 360
 caaagcaaaa cacaataacc agaggtagat ctcccgtcca tggntatcga attgaaagat 420

tcgaattgtg aatcagaaaa gctatattac gaatcgtgaa tcgaatcata tatgaatcg 479

<210> 459
 <211> 456
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 459

tgtcaaataa atcactgcat attgtgcagt ccctatcatc cttcagagga atgtagaaga 60
 gagtaggctt gcaaacaagg ttccaatttg gataaaagtg taaaactact tattaattag 120
 aattttatga atcattgttt ggaatattga agaaaaaaga caacctgaca accagctgcc 180
 ctgaaggaaac taaaatcatg gcatccaaca agaactctgc atgcttcctg catggatatac 240
 atcaataact tgaatagctg caagattgag aatctagtaa gtataaggag ctgttaacta 300
 acttgcatag ctggaagact aagctcctca ggtacatgcc atgctcgatc tttctcgaag 360
 gttgacaaaag gctctggccc agaaagcaac cgatagaagt atctgaatag tagttccaaa 420
 taactatcaa ttactgcana tggtcataaa caaaac 456

<210> 460
 <211> 470
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 460

tcagaccaaaa gcaactcana atctaggtat ctaaaacccc tcaatttagt ggattttcaa 60
 ggtttgagaa gtgaaaatga gaatgggta aacttggagc aaactctcat ctcaaacaag 120
 tctatatcat caatctaaac tcgctcaaac tggttttacg acgaaaactc taccgaatca 180
 aaatttgact cctcaacacc caattttacc ctagaaatgg ctcttggttt cactttggtc 240
 actcatattc ctcatattgca cagtctaagc tttctcataa gtcctaaatg acatttcaaa 300
 ctaggattaa ctccctttta cctccaaata ccactaaatc cagatttggc cttccaactc 360
 tcaagcctca ctctttttcc actcataaca ccacattctc actntctaac cctagggttaa 420
 ctctaccctt catctctagc agttgtccat aagcaatttc agcacataaa 470

<210> 461

<211> 248
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 461

tctacttatg tggcagggcg ggcttncttt actttctgtc tcaacgcgag ctttgaacac 60
 tgttcttcct tcccgcgatg cttcttttca tgtccgcctg agtgggctta tagcctaaac 120
 catacttccc acgatttcct tgggtattta tcacgctagt tatgccgcca ttgtctttgc 180
 ctaaaccat cccgggttca taaccgttcc ccaacataac tcgggccatc attaccgctg 240
 cgtcggac 248

<210> 462
 <211> 216
 <212> DNA
 <213> Glycine max

<400> 462

taataaaaat attatttaaa atcattaatt gagtattatg aattaatata attgttaaaa 60
 aattatagag tattagaaga caacatttgt attaaaagcg actctattat attgtagata 120
 aggtcaaagc tgttattgtg aggttcgtat atattgttag gaagttataa atttattacc 180
 tcattaagta tatttactat gaaaaaagtg actcta 216

<210> 463
 <211> 449
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 463

tctagccaaa tggacttacc ttgacttaat tcctttgata gcccttttga gccttgtttc 60
 ctttccttg ttttgaagct cactacaagc cttaagtga aaaccatgat atcactatat 120
 ctttaaggaa ttttggagct ttggaattgt tttgggaata agtgtggggg attttgtttc 180
 attggataac atgttttgtt ggccatgctt cattatatat tttgagccat acttgataca 240
 cattgcatat tgggttaaagc ttggacatgc tgaatatgat gttgtttctc aaaaggctac 300
 agaaaaaaaa atattataaa aaaaatcgaa aaagaaaaac agtaaagttg agtgaataag 360

aaaagaatga tgagactctn gggtctactc tnntatgtta aaatntatct ttacttcttt 420
 ttattttctt atggtttctt aatatgcac 449

<210> 464
 <211> 432
 <212> DNA
 <213> Glycine max

<400> 464

ttcaggataa ggatgaaaga aggcaagatt gcacgtgtat ttgtctgcat catatgtcat 60
 agctagcgat aaatcgtaga cagtagcgat gaagcgggca atgtctgttc agaaatatgt 120
 cgggttgga cagttgggtc tcatttcagg catttgtcag gctggcttgc gtctcctatt 180
 ctcttcatag gaaataatta attcccaatt agcaaagaag attaattaat tgaatgcttc 240
 agaaatttcc tttaatcttg agtcacagct ttattattat taattatatt atttcttctg 300
 tctatatatt atatatcgct gtaacgcgta tcattcattc atcaaggat ggtatctctc 360
 actgttaata gaaaactacc aacagtacag tttcttatct aaccctttga agtgcggagt 420
 acagttctta tc 432

<210> 465
 <211> 442
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 465

ngaagtactc aagggatttg cattaggtag ttcgagcaat actccttgca gaatatcaga 60
 tcttgctgca aattcagaaa gactgaagcg actcagcata catataggaa gtggggctgt 120
 gatctaagaa ggagaggttg aaagcttgag agaactgtca tcacttgagc atctcaaaat 180
 atcatggggt gtgtcagaca caaggtacgg tgatattcca atcagtttgc cttcagagtt 240
 aaaaatgttg caacttgaag gctttcctgg aaagaatttt ccagaatggt tgaatattca 300
 tagtaagcta tccagaaaat ttatgtcact atctacgata gggggaaaac ttgaaagtat 360
 ggatattctc aaatatgttt accagtacat ggggaccta ngtttcaagc atttgattct 420
 tgacacacca cnatttgga ac 442

<210> 466
 <211> 347
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 466

 tgcttggcaa ctggttactg ctgaatcctc ttgggtctct ttgataaggg ctagattttt 60
 tagagggaac cacagagttg ctttccacgt ttcttcttct atatggtata gnttatgccc 120
 cttcttagat gttataactg ataactctca atggcaaatt ggttggtaaa aacattgcgc 180
 tttggactga caaatggctc tccaacccc taagtgcatt tgttgcatat tcctgaatcc 240
 tatcatacca atttaaattc cacagtggca gactatattt ataatggtgt gtagcgcatt 300
 cctcaatctt tgcagcaatt atatccgact ttgatgaatg aaattca 347

<210> 467
 <211> 415
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 467

 ntataagcgc gggctctggga gacgaaggtc aagtggctgc gatatacgaa gatgatgttc 60
 cgagtacatt ggatttggtg cgaccatgcc ctcttaattt ccagctggga aattggcgag 120
 tggaggaacg ccccggcatt tacgcaacga gcataatgta aacctttacg gttttaaaag 180
 ctctatagtt gggcctaggc tttagagttt ttcttttgt taaggctttg tgtcttttgt 240
 tttgaattta taatacaagg atctttcttc atctgttctt acgtctctac ccattctcat 300
 tcatttgcatt gtttacttct ttttctgana atggcagatc cgatgacgag tcccctgaag 360
 gtactaatac ctgggacccg cctatcgact tcgagcaaga aatgagtcaa acgga 415

<210> 468
 <211> 329
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 468

 ntacagcaga atttagtaat gaccactaa cctagaatta aatataactt aatgccatta 60

acctagggaa ttaaaacaaa ctaaatggct gagggtgaact gaaattgttg gcaacccaaaa 120
 gtcacccccca acagccaaca agtcagccac catttgggtct cccaaaaggc tgatgcctag 180
 gttgcccaatt gggcccttat tacaacttga actaaagccc ttttagttga ttaacccaaa 240
 acatattttt ggtcagccaa ctttacaagg attgggcat tatttagaca aactaaacac 300
 tctaaaantg aaataaagtg gtgtcattt 329

<210> 469
 <211> 359
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 469

agtcggcaca ttctggccaa ttgcatgtct ccccttccct ttccctctgc tcattcgatc 60
 agctcttaaa tctaagaaca tccaagtttg aacgtgaggg gtgatcagaa ccaaaatggc 120
 caatcttagt agtcctggat ttaacctctt tttgggtgtct ctcttggtta cgttgggtcca 180
 aatccaaacc aagggtgcaat gctaccagta caaagttgga gatctagatt cctgggggat 240
 ccccatattca ccaagttcac acctctacga caaatgggcc aaatatcaca acctcaggat 300
 cggtgattcg ctctgtaagt cctcccttga tgtccattnn tttattgaga catgttaat 359

<210> 470
 <211> 454
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 470

cttggagttt ccaagtgcc aatcgctctt ttcttttagtc cagtcttctt ctggcttcaa 60
 ttcatcagtg ggctttccct ctgtgtccag catcttgga tgttcccagc ctttgatgac 120
 agctttccag gttctgctat ccagtgattt gagaaaggcc accatccttg ctttccagta 180
 ttcatagttg gtcccatcca gaattgggtg tctgttcact ggtccgcctt ctttctccat 240
 gttcatcaga atttatctcc ctgatctca ctcatgatt tcgagtgcct gctctgatac 300
 caattgaaat tctgatactg nggacagatg tcgtacagga tgtcacgaca tcacgcttca 360
 gaacatgcag atttatattg acagtgtgaa caaatataac aagttaataa cacaagagaa 420

ttgtaaccca gttcggtgaa cctcactaca tctg

454

<210> 471
<211> 426
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 471

ggctntgcan aatatatact tcattcagtg gaaggatgag aaataactaac tctntattct 60
tacaagttag aacatacaag aatgcaatca gatgttcata gaaatatttg ctgtaccttc 120
tacatcctca aatacactat ctctgattca ataaaactag agaatatcat tatacccatc 180
tctgaattga taaggaaaaa ggagttaaga gcttgattaa catcggcctc atttatttagc 240
aattttctgc aatatcaagg atcacgatga aacaaaaatc gcggccacaa tttaaaacct 300
tagcacagat gaaatctcaa gtaaacacat aacaataagc aaacacctac agcatgatta 360
acatttcaac attcaaagat cctcccatca caaccacat aaaactcata cctcacanaa 420
cacatc 426

<210> 472
<211> 453
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 472

gttgaatgca ttaaaggtaa acaaaccaaa agcaagaaat tgtgtgcata tagagctaca 60
gacgtcttgg aattgagaca tacaaacatt tgtgggccat ttcatacact ttcattggaat 120
ggcacaacat attttatatc attcatagac gattactcca gatatgcata cttgtttctt 180
atacatgaaa agtcacaatc tttggatgta ttcaaacat ttaaagttga agttgaaaat 240
caactccaca aaagaataaa gtgtgtcaga tctaaccatg gtggtgaata ctatggcaga 300
tatgacggtt caggtgaaca acatccgng ccttttgcca ggtacctaga ggaatgtgga 360
atcgtccac agtacaccat gtcgngtca cctagcatga atggtgtggc tgaaagatga 420
aatagaactc ttaacgatat ggtaagaagc atg 453

<210> 473

<211> 432
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 473

ntgagcaaat tcaaacgaac aataactttt actcggatgt cagattgagt cccgtaatat 60
 atcgaaaagc tcgaaattga atgttgaagc tctaagcaaa ttcaaacgac aaaaactttt 120
 tactcggatg tctgattgag tcccgtata tctcgaaaag ctcgaaatgtg aatgtagaag 180
 ctctgagcat attcaaacga caataacttt ttactcggat gtctgattga gtcccgtaat 240
 atatcgagat gctcgaaatg gaataccgaa gctctgagca aattcaaaca ataataactt 300
 tntactcgga tgtccgattg agtcccgtaa tataatctgaa cgctcganat tgaatgtcga 360
 agctctgagc aaattcaaac gacaataaca ttttactcgg atgtctgatt gagtcccgtg 420
 tataatcttga cg 432

<210> 474
 <211> 368
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 474

tccaccggtt gtgattgcga gataatattc gtggaggagg aataaggaat cgtatgaaga 60
 cagtacaagt ggaggtttca atctcttctc cgtctctctg acttttggga attctatcgg 120
 agtagtcgga tgaataattg aaagaatttc tgggaaccgc tagagatggt gttatcgtg 180
 gctgaagaca cgtgagcccg cttagaggta agggatgagt ttatcgcaaa tgggattaga 240
 atgaacatgt gtanggatcc ttagagaact aaatttgggt taatttgcga tggttattga 300
 aatataattt ctctttatga ttataaatat aatattaatg gggcttatgt accaatgatg 360
 ttctgatg 368

<210> 475
 <211> 409
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 475

ntgagccaaa atcctgactc accatanacc ttgaccagg gtgagaatgt caatccttac 60
 cctcggaagc gaaaagaata gaagggaat ttccaatcaa agaaaaggaa agaaggaaga 120
 tttccaatca aagagaaagc aaaaaaagaa aagaaggaaa attcccaatc aaagagtggg 180
 agaaagcaaa aagaaaagaa agaaaattcc caatcaaaga atgggagaaa gtaaaaaagg 240
 aagaagaaga aggaaagata gtcctgatc agggatcgaa agaaaacaga agaatgtgc 300
 agaaaggtct ttggaccgga caatatctga ataatacaga gttgtcacca aatgaacaaa 360
 aagaaggaaa ggaaaccacg acctanaatg gtcttctccc tttgattac 409

<210> 476
 <211> 434
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 476

tgagtgtcca catgggtgca tgcgatgatca attntgtata agtttctaaa tcatcgttgt 60
 tatatacgtg tcatggaaat aatgtggggc acttcctttt atccctgaac cgctggccaa 120
 agcaaatacc ctgacatacg tcatgtcttg ctttctttaa agcctctttg ggacaaaacc 180
 tcaatccttc agccctaagc ctcgacccaa ggtaggaatt tttaccctta tctcgaaaa 240
 aaagaagaac aggaggatct aaaaaaacg agaggaagaa aaagtttcat ttacttttaa 300
 gttatgaatg tgccttacaa ggaaaataaa aagagaaaat cccaatcaa agattggagg 360
 atagcaaaag aaaaagaaaa agaacaattc ccgatcaaag atcggaagaa agcataagaa 420
 aaatatacag aaag 434

<210> 477
 <211> 426
 <212> DNA
 <213> Glycine max

<400> 477

tgctcgtaa gcctgtgctt tcttcttgag tggttgcgct aagctcggct tgccgcacta 60
 agcgctaatac tttctttgtc ttaaaaaatt gtggaattag gcttagcgag caggcttcct 120
 aagcctattc tgcagaaaaa aagattttat gtgttcttgc gctaagagca tggctatcac 180

gcttagctta tgagtaaaat ttcataaggc gcactaagtg catctgctgc gctaagcgcc 240
 caatcttaat tctagattta ttttttgctt ttcttttggga ataattcttg tctagtcttg 300
 gcttttgatt cttttgtttt tcagatggct tcatgaaaga ggaagacaac agctgcagta 360
 ccccgaggccg gatatgacat atctagattt acatcccaag aggcattgtga ctgctacaca 420
 tataat 426

<210> 478
 <211> 514
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 478

atgtctctgg gcccttggac ctgtanance tcttttgnnc ccttgaaccc ccgcggtat 60
 ttaaaaaaac ccgcccctga caagaaagca ctgtggagat gcttaccacc tcttatgact 120
 ggaaagcggg ttctaatacgc tcctctgcgg tctccacata aggcataatag gaagggcagc 180
 tcaccaagat gtcttactcg cctgatacga tgaccagatg cccttncact acaaatatca 240
 acttttgggtg gagcggagag ggaacaacta cactgagtg gattcacgga cgccccaaca 300
 gacagctgta gaggagggtta atatccatta ttggaaagt cacttgacag gtgggagggc 360
 ctattcagtc tgggagaact attctcttcc caaaccttct cggggtggtc tcctggcaca 420
 acccactttt gaccatattt gtatgagaac aaagtcgctg gtcactcttt gggttaaccc 480
 atgaattgat taatgattgt gaatatatcg agag 514

<210> 479
 <211> 456
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 479

ctattacgtg aactatgaa actaagctnt aacanatgtc ttcacaaata atcatcacac 60
 agcagaaacc tagcaagact acccatcata tctcccaaaa ccccataccc acaaaaatta 120
 aaggagaaag aagtccaccc aaacctgaat tttcgaagtc ccactcgtag ccacgcactt 180
 cacgaccccg aaaatgcctt cctttcgcca tttggggcag aaatgatggc caaagggttga 240

agcttttgctt ggagcttcaa tggagaatga agaagaagaa aatggcaacg tgagagagaa 300
 agagagcttt ctgaaaagtg tgggggctga gtgaagagag agaaaagctt tttggtnta 360
 aataaatggg gtntctcttt ttctattatt ntatttaagc aaatgccaca tgtctncatt 420
 tgagtggagc aaaaagggcc cacttttcct tttgac 456

<210> 480
 <211> 519
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 480

tcccaccgtt ttttgaagca tgctattagg acctatgaaa ctaagctgtg cttanaggaa 60
 aacttanata aagataactn tcatttcaca tgtaggtgt agctggtgta ttcggcggct 120
 cctgtataag gataaggggc gaaaagtaat caaatagaaa ttaaactaag aaggcaaagt 180
 gtgaatgtgt taaattgtca ataatctcac ttcaaccatt caacacagat acataagatt 240
 atttcagcct aatcgataat ttttaattta aatataagta tataattgta ttaaattattc 300
 aaaaagaaaa ttttgttaca caatcatatt acaatcaaat aaaattgtgt ctccgggtag 360
 agatacatat agtccctact caaaaaatcc catttcaaaa gtaaataagg caaattacta 420
 atgttgaaaa gttttgtaat catgaaatat gtcgcttcct tactagacct gacagggtcat 480
 ttaatatatt attatcaagn attgtttacg cctatgaga 519

<210> 481
 <211> 245
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 481

tgagctgcan atgttgcttc accatctgtc aactgtcacc tgtacaactc tggcacgtgc 60
 ccaagtgaaa tgcacatgtg ccttatctgt aactacacat ggcttcttta acgtcatcac 120
 gagaaagagc atgtacgtgt agggcttccc aattaacaat ggaagatgta ggaggcatga 180
 catataatcc antgtgtgat ggaggagag gggttgggaa agaaatgagg ggtggtggta 240
 gggggg 245

<210> 482
 <211> 451
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 482

ctgcccattt aagttttgcc agcgaanaaa tcgaagtggg tctgagaaga ggcaaatttg 60
 attatcctgc tttgatgaat aggaagcttg gggcaaattg agagaataag aaggagggaa 120
 aatcagtcct tctcattacc caccacccta ccagccatga acgcctaata atccacaaag 180
 gccatcccca aatcagccaa aaatccaccc gatgcacatc caagaccaa taccaccctt 240
 aataccaatc aaaacaccaa ctagggaagg aattttccag aaaagaagcc tgtagaattc 300
 accccaattc caatgccata tggtgactta ctccctacc tgctcgacaa tgcaatggta 360
 gttataagcc caacaaaaat ttctcaacct ctgtttccca gaggatacaa ccncaacgtg 420
 acatgttctt atcatggggg anngtttggg c 451

<210> 483
 <211> 449
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 483

tctataaaag gttcgttctt aattttctta caattgcata acctctcaat gagctggtga 60
 agaagaatgt ggcatttacc tgnngtgaaa aacaagagca agcctttgct ttgctccaag 120
 aaaagcttac taaggcacct gttctagctc ttctgattt ttctaaaact tttgataata 180
 ttagggactt gtatgcttta gatgaacatt tctctcccat ttacgaaagt tgtgggaaaa 240
 aggcccaaaa tggattctat ttggctaagg ggtatttgtt caaagaggga aagctttgca 300
 taccccaagg atccattagg aaattacttg ttaaagatag ccatgagggt gggctcatgg 360
 gccacttttg gatagacaag acgctcgtct tactcaaaga anagttttat tggccccata 420
 tgaagaaaaa tgctcttaag cattgcact 449

<210> 484
 <211> 405
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 484

actctcttat cgaatggtgt gccacaataa catcaacatc gatttgatga ttgcatgctc 60
ccttttctga tcttataggc acgttacagg gacgaaatca ttatttaaact atttaagtgc 120
ccaaagattt tgattntaag agcaaaaataa aataaaaaaa atactgaatc atgtggagtt 180
tattgaactt catatgcccg agtagagtgt catgcatggt aatttaaact gaatcaaact 240
aaactgtacc ttacacatta gtaatgtaat tagtaagact agaagcatct taatgagggtg 300
gcagaatnta attaattatt tagaaacatc ctaatgaggt ggcagaattt aattaattat 360
ttgatttcaa tatttttcat actaatttct tctnctttat ccctt 405

<210> 485

<211> 398

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 485

tgaagagagt tcttccatgc cagcacgcgc ccaacgtgta tgtagctcgc taagcgagat 60
ggttgctctt tctgcgctaa gcgcgagatt gccgctaagc caaatctcac ttactcgcac 120
ttagcgcgag aatggcatta aacgcgcctt catggacagg aagccctttc ttaagcctga 180
cttacagaaa atgaagggga gggctggaag agagcgcgtga atagccgtca gagtttgaag 240
agtgaatac acaaaggcaa ataacagagc anaggagcca agttttgatc ttttaggaag 300
atgtgtgagt ctttgagtga ttgtgagatt cctagagggtg gaggagacat cctcactcct 360
ttttagcaaa gcaatntctc ttaattcctc ttctttca 398

<210> 486

<211> 382

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 486

tcattgatca attctgaagt tcacaattgt cataagttca tgaaaggagc aaaatgttgg 60
atcaagtggc ctcagaataa ttaagaaggg gggttgaatt aattattaat gagcctttac 120

taatcaaaaa cttatccttc ttaatgttac tagattcaat taggccttta ctactaagtt 180
aagaaagtaa agaacagaaa tagaaactta atcaaatgta aaagcaataa ttaaagtgca 240
cagcggaaat taaagagtat aggggaagaac aagacaaacg caagaattnt atactgggtc 300
ggcaaaaactc atgcctacat ccaatcccca agcaacctgc ggttcttgag atttctttca 360
accttgtaaa atcctttaca ag 382

<210> 487
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 487

nttggcttta gaattaatca tcaaaagtct catttcattg tgtctaagaa tattccgaga 60
aggaaaactc anaaatttgc tatattttgg gtttccaata tatactcatg atattggtaa 120
atatttgggt ttacctataa ttagtggaag agttaaaaaa aaccacttct cgtttattct 180
ggataaagta aatgatcgt gagctgggtg gaaatcgaag cttctcaata gacttggtcg 240
ggttacactt tgcaaatatg tcttcaattt tatccttaca tatgtcatgc aaaacatgtg 300
gctccttcaa ggcatthgtg attcccttga tattgctact agacaattca tctgcggatc 360
aacttcatct cattgggtga gttggaagac tnatcatc 399

<210> 488
<211> 396
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 488

tcttggcaat cctcattcca gcgatcagtt tggtnnttgc gtaagagttt gaacaacggc 60
tcacaaatgg cggtgagctg cgatatgaat ctggcaatat aattcaagcg tcccaggaaa 120
cctcggactt gcctctctgt acggngttct ggcattctcaa ggatagcctt caccttttcg 180
gggtctacct ctatcccttt ctggcttaca acgaaaccaa gcaatttccc tgatttgacc 240
ccaaagggtac acttagcggg gttcaacctt aattgatatt tcttaagcct ttcgaacaac 300
ttccgctggt tgacaagggtg ttcttctctg gatttagatt tagcaattat gtcgtccacg 360

tagacctga tctcttgatg catcatatca tggaac

396

<210> 489
<211> 383
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 489

tcatggtgaa tcaaaggatg ttcanaggatg ttttgatgat aacaatgatg ataacaaaag 60
atgatgacaa aggtgatgac aaaaagctca aagatcaatc aagaacaatt caagagttca 120
agataagaat caagaagaat tcaagactca ataagaaagt ctagagacaa gaatcaagat 180
tcaaggttca agatctcaag aatcaagaac aagattcaag actcaagatt caagaatgaa 240
gagaagactc aatcaagata agtattaaaa agtttttcaa aactttgaa agcacatgag 300
tttttgacaa aaacttttac cagagttttt actctctggt aatcgattac catatagttg 360
taatcgatta ccagtagcaa aat 383

<210> 490
<211> 346
<212> DNA
<213> Glycine max

<400> 490

tctacttatg tggtagggcg ggcttccctc actttcttgt ctccaacgag agctttgacc 60
actgttcttc ctccccgaga tgcttctttt catgtccgcc tgagtgggct tatagcctaa 120
accatacttc ccacgatttc cttgggtatt tatcaggcta gttatgccgc cgttgtcttt 180
gcctaaacct atccccgggt cacaaccgtt cccaacata actcgggcca tcattactgc 240
tgcaacggac agacaagggt gccagagag ggagtccacg gaggaatgc tgaccacctc 300
ataagactgg aaagcgggtt ctaacgattc ttctgagggt tccaca 346

<210> 491
<211> 391
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 491

tgaaggtaaa aggtttacac agtttcatgt taaaagttac tgacatcctg ctgctaatag 60
acattagacg tagctntact cttagtaaga atattagaac aacacagcaa aaaaaacact 120
ttctatgcat tgagcaaagt tattcaaaaa ataattatgg attagaacta agttttcaca 180
aatcttaagc aagcatcagt aacatcttta cctgcagcac tagaaagaac ccaattgtca 240
tcataaggaa cttgccacac agcatagcca agtaacttat tttcccttgc ataagaaacc 300
ttcatntga caacctcaac atcatcataa cctatccaag tcgatccatt ggagaagtaa 360
ttaactacat aagtagcatt gtacttgaca t 391

<210> 492
<211> 454
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 492

ctcagcttan atatgttttc cctcanatga nnactaattt agttntaacc catcaattaa 60
aaatttgctt tctacaaatc ctcaaattca ctttgaggca ctgaaatcaa cagttttacga 120
atttgagtga tttaaaacat aaattatgct catgtaacct ttcatgaata aagactaaat 180
tattcattca aaaagaataa atactgaaac agtatgaagt taatgaattc attgactaag 240
taaataataga aggacaaatt aaatacaaaa attatattat atgcaagtaa tcttttaatt 300
ttaatataaa attaattagc acacactatc acttttgaaa atgatatata tatatatata 360
tatatatata tatattaatt ctaacattag tattcgtata gtatacacgg attgttaata 420
ataacaagtg aacaacaata acagtaataa taaa 454

<210> 493
<211> 386
<212> DNA
<213> Glycine max

<400> 493

tgcaacttcag gacgtgcaat gctggtaaag aattgtggca ctgaactact tgagttgtcc 60
tcaagggtaa ccctttgttg gtgttcatca tccatggtaa tggcctctgg taagtttgaa 120
gctgattccc ttgataatgg tgagtcagat gatgaagaat cagaaaaatg agtttcttcc 180

tcaagaatgg atgctactgt cctgtcttgc agtaattttc ttctcctttt ggctctgttc 240
 cttcttaatg tagcttcaat ctccaagtcc aaaggaacta aattgcctgt gggagatcta 300
 tgcattttaa acactaacag aaacaacagt tatccagttc aagaggaaaa aaaatatgaa 360
 ttaaaagcaa atattcacag ttaatc 386

<210> 494
 <211> 443
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 494

tagccgaatt catattgaat tgaagttagc ttagctcaac cttggccagc ttagtggacc 60
 aaatcaacct tatgcaaggg ttgggcgcta agcacttgag actcacaact tagcgcatga 120
 accgagatgc gcttagcgta aggcttgcgc ttagcgaaag gactattttt cagagaaaag 180
 ttttctgtta ttttctagtc ctttttccaa gaaattgaaa cttttatgtt aaacattcaa 240
 agataggttg atatactcct atgtacagat ccgacagcaa gttccaaatg attaaatgca 300
 tgaaaaacaa agataacaaa atttaaaact gggttgcctc ccaggaagcg cttctttaac 360
 gttattagct tgacgctntt accttactgg atgatcttat gttttggttc ttactttcag 420
 aacctcttga cctccttcca tta 443

<210> 495
 <211> 369
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 495

nttagcttct ctaggaatct tctcaaggaa gtttctcaag gaagcttctt aatgaggtga 60
 gcttagttat taggtgtgtg tagctaagtc tagcttctca aggaagcttc tcaaagaagc 120
 ttctcaagga agtttcttaa ggaagcttct caaggaagtt tctcatggaa gtttctcaag 180
 aaattttctc aaggatgcta cctaggctat aaataaaagc atgtgtaaca cttgttgcaa 240
 ctttgatgaa tgagagtctt gtgagacaca cttcaaagtt caacttctct ccctctttta 300
 caccttcaat ttcatgctcc ccctctctc tttctctctc tttttctttt tctccattga 360

agcttccta

369

<210> 496
<211> 372
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 496

nggatgaagt tegaaatctc ttcctttcttc atctctaaat ttcattccacc accacttgca 60
tcaccaagga gagctccaaa actgtgcttt cctcaagctc ttactctact ttctcttaaa 120
ctctctacta tgagtgtttt taagtgtgtt aaacccaaat aatccttggt gtatttatag 180
ggtaaagtgt aggcataagt agtaaataag accaatgagt gtttaaggatc atataggtct 240
ttaggttaca aattaattgt tcttatcttt taattntatt tttttctttt cttttattaa 300
ttagatatcc tagatgcttc atgggttatt agagtagaga ttggaatgta tgtatacatg 360
atcttgatga tg 372

<210> 497
<211> 387
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 497

tttcgcanag cttacggtaa aatctaggac atagccatgg caaagggtcta cacagaggcc 60
attgcctccc tcgcccagta ttatgactag ctgttgaggt gcttcacctt tggggacttc 120
cagctatcac ccatgggtgga agaatttgaa gagatcctag gatgccctct atggggaagg 180
aagccatacc tcttctcagg attctatccc tctttagcta gaatttctaa gatagtccaa 240
atctcggcgt aggaattaga ccacagaaa caagtcgaaa atgggggtggt tggagtaccg 300
agaaaatggt tagaggcaaa agcaagaatc ttggcaggta gaggcaaagt ggccccgttc 360
atagacatcc tcgcactttt gatcttc 387

<210> 498
<211> 446
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 498

tcagcatctt aaacagaggc taaaaatctg aagtaaagac aacatatgag acctgtgcag 60
catagtcaaa cagcttcaat agaaactaaa tgacttggag aattctatgc cagttcaacc 120
ttctgaacaa caagtgaagg acctcaagaa aaccaagct gacctttggg aaaaagctac 180
tatgcaggag tctattgtga ggcagaaatc aagatgtaga cggatcatag agggggacag 240
caacacatcc tattttcata gagttattaa ttgaggagg aagagaaatg ctctgagggg 300
gttgacagatt ggtgacacct gtgtggaaaa tcctaacatt atanaagctg aaacctttca 360
tcattttaga acaggttcaa tgagcctcac ttgaccagac ctaacttga tgggggtttc 420
atttaaagtc tgacttattc tcacag 446

<210> 499
<211> 402
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 499

tcaacatgaa gcaatcagct cgctgggca agcatgttgc ttctaaacta agccaccagc 60
ttgccgggtg agctgggagg caagatcctc ccctattttg gctataaaag ggtgtgggag 120
gctaagggga aggggttcag cacccttggg cacttgcagc agacaaggaa agtgtcggtg 180
aactgtatg caaaaagtac gacattggca atgagaagtg ggctcaattt tgtcagaccc 240
gcagagaccc ttcgtgggag gcaacgtttt tattttcatt gttntaaact ctaaattcac 300
ttagtataat acattgtaat gataactttc aataatgggt aacttttaca ggatatgoga 360
aaaaaggcat aagccatcta aaaacaaaac actgtccctc ac 402

<210> 500
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 500

tgcttgtggn gcttctatgg aggctggatc ttgagcttc aatgaggtcc ttcaatggtg 60
attttcgacc atggagatgt agcagaaggc aaaggagaag aggagagagg aggcaccatc 120

cacaagggaa taagccaagg aagaaggagc ttcgccacca agatgagcat tggataagaa 180
gcttggagat gatgcttcaa tggaggaaaa gaaagagga gagaaagaga gagtgaggag 240
cacgaaattg aaggaagaaa aagggagaga agttgaactt tgagttgtgt ttcacaagac 300
tctcatccat caaagttaca acaagtgtta cacatgtttc tatttataga ctangtagct 360
tccttgagaa gctntcttga gaaaacttcc ttgagaagct tctttgagaa aacttccttg 420
agaagctaga gcttagctac acacacccc 449

<210> 501
<211> 373
<212> DNA
<213> Glycine max

<400> 501

tgagtttaac gatgacaaat attcaagaag caaacattaa gttatccgaa ccaagctatc 60
tagtattgat tgcttttaggt gttctttata acatacgact aaacttataa gaatggggaa 120
gatgtggagc agtagcctat ggtagaaatt ttccagacat agattctgcy taatcaaaaag 180
caaaactact agaaagagag tggctctaatt ttagatggaa aggaacacac gttttatgag 240
tgaaactgaa aatataacaa gctatatatg atcagaagca tgatgttagg tatctttggt 300
aaagggacat atattagaag ctacgcatga ctaagggatc atccaaagta ttctaattca 360
aaagtcatga gaa 373

<210> 502
<211> 320
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 502

ntngagatca aactntacca ctggtaatcg attacaggaa actggtaatc gattaccaga 60
gagtaaatac tctggtaact tagaaaaatt tggaaaaact tttcttgtaa aacaaaattg 120
tgctatgttt gggttttgaa aaatcttttt caatacttcc cttgcgaagt cttgacttgg 180
tgctttttgg tttcttctct tgaatcttga atcttcttga tgacttttct ttaatcttga 240
tcttgaactt gttgactcaa tcatgacatt attcttttgg catttttgaa atcatcaaaa 300

ctacttgaat tattcttgat

320

<210> 503
<211> 430
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 503

tgggtggtaa aaananaaat ataagaaaaa ggtgtatgtg tattttggga ggagccctgc 60
cgattcctaa atcctctttt cagaggaaaa cgtaaagaa atgctactgg aataagataa 120
gataaaaaat ggagatgaga gcatgctcca ctgattagat tctgtaagcg ttagctacta 180
aacttcatca acaacatggt aagtaattat ggcaacagtg atcgaaccaa aggaatatgc 240
agacaagata gatagagaga aagggaaaaa aagtcaagtt aagtgagtgt catcgttgaa 300
aatcaatata gatgccaggg acggctctgt aaattcagta gcccacctta caaataagtg 360
ccctaattat ctgaacactc ttctcaaata atacattcac aaattaaaca gttccattag 420
taaaaatatg 430

<210> 504
<211> 463
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 504

tattggccgt tggatgaaac tccacacagt gcagttctat gttcacttgt agaggatctg 60
aatcccagtt taccagttgg ttgttgccgt taaccaacaa atcttatttt atgttctctn 120
taatatttct ttntaattga agatgcttct tttaaccaca gaaattgtat atttttcatg 180
caagtataac agttaaataga atagtagtta gtttaattgt atttcgaatt tttttttata 240
attaaattat aagtgatttt ttagtttcta tggaaaaaat taagttgggt aagtttttta 300
ttttaaaaat taactaaatt tattcatnt gtcaaattnt tacttaaaaa ttntcaaatt 360
tttaatttta tctacatatt taggagttta tgaataaatt gagttagaaa ataattntaa 420
aaattatang taactttatt agaaaaatta gtaaataatta atg 463

<210> 505

<211> 406
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 505

tcttcacctt tgctcagagc tggtaattga acttattcta aaattntgtc ccttggtcca 60
 aaattatggg tgaattttca ttagattatt ttttcgcttc acataaaatg attctttgat 120
 catgtgagaa gagaatgaag cctcagttag gacagttgat taaattaaga atagactaat 180
 aattacaggt agaggaagac caagaaagac tttggattct gctattatta gaattggttt 240
 agattttaat ggcttctatg aaaaatgaaa aattgttttt taacagaata caatggcatt 300
 gtttgattga tataggttgt agatgacctc agtgggaaaa aaaactntgg ttatcactgc 360
 tatgtataaa tcattgatat tatttggtag caactcttat ggtaaa 406

<210> 506
 <211> 460
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 506

tagcccctaa cgaaagatga accgtgtcaa gttgtacact atgtggagat cattgtagga 60
 gtagaagggt ccgaggagtt agagttggac ctgagaacca atgacgacaa tcgggttaaa 120
 ccaattgaag aaacatgtaa ctttcagctc gacattaaag aggaataggt tacttgactt 180
 gggaaccaac tctcaatgga atataagaat gacttacaac aaattatctg agcacatgcc 240
 gacctgtttg catggttcga ggtcaacatg ctaggcatag atccgacctt ccattgccat 300
 aaattatcca taatttaaga tgccaaatta tatctcagag gaagagaaag attgngaaag 360
 aaagggtgta ggtagtgcgg caagaggttt ccaaactggt ggcctaatta tcanagaggt 420
 agagtacacc acatggctat canatctatt cctagtaaaa 460

<210> 507
 <211> 459
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 507

ggcatagtat gattgcactt ctggctctat ccatcatttc ttatttctcc tttgaactta 240
gagattcaga catcctttct tcttctttaa gagcttctac acagccatgc tgaatcaaga 300
ttgcttncat cttgatcttc cataaccga agtcattttc cctgagaact ctcatatcat 360
act 363

<210> 510
<211> 418
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 510

tgattaccct ggtgacactn tacttccagg aatgaaaatt ggagtgaact tcaaaaccgg 60
tcaacatcga gctctaagat catggagatc ctttacggat cctactccag gtaatttttc 120
acttggtggt gatactcgtg gccttcctca attagttatt acaaatgaga atactaatag 180
taatgacata gcttatagac cagggatcatg gaatgggtctt agtatcacgg ngcttcctgg 240
agaaataact gaccaattaa caaaatccct ttttgttatg aatcaagatg aggtcttcta 300
tgagattcag ctcttgaata gttcaactaa actcatgaga agcagaatgc ttccagaagg 360
gtatcaagta cgttttatat ggtcagatga aaaaaaatat gggattctca aattccta 418

<210> 511
<211> 435
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 511

tagaagcaaa tgctggcatg gtatcgaga ttcgtgtagc aagaccacct gctgaaggac 60
ggngaaggaa tcaattgctt ccacgttatt ggctaggat tactgatcaa gagttgcaac 120
aaatatcagg agagtatcca acattgtaat ctgtcctata aattattatg aatcactgac 180
aggatactta ctggttgata cacttattat tattttgata gctaattctt accttagtat 240
ttccttaacc atgattcatg atatgttcaa gttcaaattc tacaatcgtg ccactctttg 300
aaaagatgct tagtgcaagt gatgctggtc gaattggctg cttgggttta ccaaagcat 360
gtgctgaagt aatttatctt aactcatctg ttgaactggc atttactgtt gtcattttat 420

attaactaac aattc

435

<210> 512
<211> 395
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 512

tgtgaatnta tagtgtttgg tgactaattg tcacaaaaaa gcaaagtaaa gcccaaagaa 60
gcaaaattaa agatctaaaa ttactcgctc agcatttctc aggtgctcag cgcaacgcag 120
atgcttagcg gacaacgcac gcttaacgcc agaaagtatg aagacgtctg aatcatgaat 180
atgtgcttag cgcgagtcac tcgctaagcg cgagattact atcatactcg ctaagcatga 240
aattgcactt agcgtgaagg ttacgtaaaa atcaaactga actacaccta taaaagaagg 300
agagagaaaa agaaaaaaaa tacacttaaa attcaagaga atacaattcc ttacagaagg 360
caaaggtcga aagcaggaga agcaaccatt cggag 395

<210> 513
<211> 412
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 513

tggtaaagaa aatgatggca tgacttttat ccaatcttat tatgttgaaa agctattgaa 60
gaagtttaat tattttgatg cgaaacatgc tcttactcct tatgactcat ccatcaagtt 120
aaagaaaaat ttgagtaaat gaatttcttt acataaatat tctcaaagta tcggttcttt 180
gttgcatctg acaaacttct ctatgcctgt ctgatattgc acatgcagtt ggtagattgg 240
aaagtaattg agggatttag tgatataaaa ttgaagttct gattntgatg aaataaaaaat 300
gagaagtggg tatgtctttg ctttagctag ttgtgcagta tcatgaaaat ctactagaca 360
agttattatt tcacatgana gcaaaaatta ttgctttaaa tactgctact ag 412

<210> 514
<211> 320
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 514

cttgtgtggg acacccattg tgagtgtagt ttccaaaccc ttatagaaaa gttgacgatg 60
actcctgtgt tagttttgct taaccaaga gaaccctttg aggtgtattg tgatgcatca 120
aagatggggtt taggaggagt gttgatgcaa aatggacaag tagtggttta tgcttctaga 180
caactcaaga ctcatgagag gaattatcct acccatgatac tggagttagc tgctgtagnag 240
tttgccttta acgcgtggag gcattaccta ttgcactcca gtttgaagtg tttagcgatc 300
ataaaagcct taagtatttg 320

<210> 515
<211> 371
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 515

taggaaccca nacttgtagc ttcaatgcaa ggaaacgtgc ttatggctag gaatccaaaa 60
atttggtttt agaattagaa aagaatgaaa atagggactt gtttgtaaga atttgggctg 120
ccccatgatt ggtactttgc acctaaataa catgggaaat gattttcaat gctgtgtaga 180
tatatgtgta aatatgaagg gcatgaaatt ctttgcaaag gatgaaggaa tattgaggtc 240
acttcctaaa tgaatgtatg atagcatggg attccctttt gaatgcaagt atgtgcataa 300
tgtaaataat cttgccaata ggcataagtg tgagtgaac aatgaaaagt tgtatgggat 360
atatatcttg a 371

<210> 516
<211> 455
<212> DNA
<213> Glycine max

<400> 516

tagcccccta acgaaagatg aaccgtgtca agttgtacac tatgtggaga tcattgtagg 60
agtagaaggg tccgaggagt tagagttgga cctgagaacc aatgacgaca atcggggttaa 120
accaattgaa gaaacatgta actttcagct cgacattaaa gaggaatagg ttacttgact 180
tggaaccaa ctctcaatgg aatataagaa tgacttacia caaattatct gagcacatgc 240

cgacctgttt gcatggttcg aggtcaacat gctaggcata gatccgacct tccattgcca 300
 taaattatcc ataatttaag atgccaaatt atatctcaga ggaagagaaa gattgggaaa 360
 gaaagggtgtt aggtagtgcg gcaagagggt tcaaactggg ggctaattat caaagagtag 420
 agtcaccaca tggatatcaat ctattctaga aaaaa 455

<210> 517
 <211> 461
 <212> DNA
 <213> Glycine max

<400> 517

tgcttgtgga gcttctatgg aggctggatc tttgatcttc aatgagggtcc tttaatggtg 60
 attttccacc atggagatgc agcggaagac aaaggagaag aggtaagagg cggcgccatc 120
 cactatggaa taagccttgg aagaaggagc ttcaccacca agatgagcct tggataagaa 180
 gcttggagag gatgcttcaa tggagagaaa gagagggggg ggagcacgaa attgaaggaa 240
 gaaaaagggg gagaagttga actttgagtt gtgtctcaca agactctcat tcatcaaagt 300
 tacaacaagt gttacacatg cttctattta tagactaggt agcttccttg agaagctttc 360
 ttgagaaaac ttccttgaga agcttctttg agaaaacttc cttgagaagc tagagcttag 420
 ctacacacac ccctctaata actaagctca cttctttgat a 461

<210> 518
 <211> 460
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 518

tggagaggat gcttcaatgg agganaagan naaaggagag aaagagagag gggggaggac 60
 gaaattgaag gaagaaaaag ggagagaagt tgaactttga gttgtgtctc acaagactct 120
 cattcatcaa agttacaaca agtgttacac atgcttctat ttatagacta ggtagctttc 180
 ttgagaagct ttcttgagaa aacttccttg agaagcttct ttgagaaaac ttccttgaga 240
 agcttctttg agaaaacttc cttgagaagc tagagcttag ctacacacac ccctctcata 300
 actaagctca cctctcttgag aagcttcctt aagaagattc ttaaagaagc tagagcttag 360

ctacacatac ctctctaata gctaagctca cctccttgag atgagaagct agagcttagc 420
tacacacccc gtataatagc taagctcaca tgaaaataac 460

<210> 519
<211> 405
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 519

ntgagccaaa atcctgactc accataaacc ttgacccagg gtgagaatgt caatccttac 60
cctcggaagc aaaaaaagaa tagaggggaa atttccaatc aaagaaaaag agaaggaaaa 120
tttccaatga aagcaaaaaa agaaaagaag gaaaattccc caatcaaaga gtgggagaaa 180
gcaaaaaaag aaaagaagga aaattcccca atcaaagagt gggagaaagc aaaaagaaaa 240
gaaaggaaaa ttccaatca aagaatgaga gaaagtaaaa aaggagaag aagaaggaaa 300
gaaagctcct gatcagggat cgaaggaaaa acagaagaaa tgtgcagaga ggtctttgga 360
ccggacaata tatgaacaat acagaattgt caccaaata acaaa 405

<210> 520
<211> 452
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 520

acaattatct aatcattcca atccactcaa atcatcacgt tgcatttca aatcattctc 60
aaacactcat ttcatacaaa acaatccact gcatatcggt ttcaatcagt tctctgttca 120
aacaagcttt tttgtacatg caaacaactc atagtactaa aatttaaaga acggaaacat 180
aaaaactgaa atttaaatga ctgaacataa atcataaaat aattcaagta aactaaaatg 240
ttcaaaatgc acaaatttaa atgtcctgct tctgtgggtg ctcttgtgca tgcttattaa 300
gatccaacac ctgagcaact ggtaaatcct gagaggtagg tttctctaac tcacatgttg 360
gtgaagatgg tatggcatca tcaggatatag gtgctgggga tggctctggg atctggtctg 420
tggaagtctn cttctcttga gccatgtgta ca 452

<210> 521

<211> 391
 <212> DNA
 <213> Glycine max

<400> 521

tcattctcta tcttgagact cttgtgttat taattactgt ataaacctta gggttttctc 60
 attcctatct tctgcaaatt ctctacaag gctagaaata tttctgagca aatataccag 120
 attttgatgt tgattctttt agcatgcaac tctatattaa tgctgctgca ttgtaatgat 180
 cacatttgca acttactaaa atgcggcgca gcttaagcat ttgaatgctt tttgaatcac 240
 tgtatttggc gtatttggtt tggccgtag ccaattcaat gcgtggatat atatcgaca 300
 cctacaccgg agatatatat cagttatggc tttgtgttca agttttctca atcttcatct 360
 tctgggtttt gtttggttga cgatagaagc g 391

<210> 522
 <211> 418
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 522

tcatttgcaa ttcagtgtaa tcagtgccag ccatattcga atgaattgga tactttgaac 60
 cagtttatgt acaaaagaaa aaaaaagttt aatttttatg tatcaagtat acgaaatttt 120
 ataccatcaa ccaattaaaa gttatatata attggaataa ctattataat aattatcata 180
 aaaatcaata aatttatcat tcatatattt caataccaat agccactata taattactgc 240
 atgctcaatg agaattaaat tgacatgtac ctctgatcgt atagtgtgta tgaatctgcc 300
 atctcttctg attgaagtag tgctaaccac gttaagccct tctttaagca gaatgtccaa 360
 cactcttgac atgggaaaca aacttttccc anagctgtaa ctgcacataa tctcaagt 418

<210> 523
 <211> 393
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 523

tcaggctgtt caattgctnt aaattgttgc atagaagggc aaaggctctgt gtgggtggtcg 60

gcagaggagc acaaaccaca gagtctggcg acaggtgcag attttttatt catggccagt 120
 tgggttacca ggtaaccaa tgcatttagt ttaccttcaa gcttcttagt ctcacctgat 180
 gaattcgtgg ctacttcatg cactcctcta atgacaatag catcacttct ggcaactaat 240
 gtgtgggagt ttgaagccat cttcttaatt aaatttcttg cttcagcagg ggtcatgtct 300
 ccaagggctc caccactagc agcatctatc atactttctc ccatgttgct gagtccttca 360
 taaaaatatt ggagaagaag ctgctttgaa atc 393

<210> 524
 <211> 392
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 524

tctaaactnt aaacaaaaat gaagctaaac tntaaacaag aatgaagctt cgataccact 60
 tgtagacaa gtggcctcag atatcttaag aaggggggtt gaattaagat attacaaact 120
 attttcccaa ttaaaattct actttgattt taatgcaagt tcaaagttcc cttaaagatt 180
 aatttctaaa tgatgattca aaataaccaa actgaatgta aaagtaaagc aacaataaat 240
 aaaagagttt aaggggaagag agagtgcaaa ctcagtttta tactggttcg gccacaccct 300
 tgtgcctacg tccagtcccc aagcaaccca cttgagagtt ccactaactt gcaaaaaccc 360
 tttacaagtt ctgaaccaca caaggacaac cc 392

<210> 525
 <211> 209
 <212> DNA
 <213> Glycine max
 <400> 525

ggagaatttg taagacttaa ttcaccctc tcttaagtta ttgaggtcac ttggcgcgca 60
 cacagatcaa gaataaagct aagtcttact ctatctttgt taaaagagtc tcttagtgat 120
 tggaagaagt tggcctcaca acattgttct taaattgatt ataaaaaggt tctagaatac 180
 ttttaacaat ttttggataa gacattttt 209

<210> 526
 <211> 431

<212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 526

tgagcatagc agctntgtta attaccttgt gagagtgaac accaaggcca cccaaatccg 60
 taggcaaaca aacctttttc caaaaaaaca taacaagttt tgatcagtta tccctttccc 120
 ctgtccagat gaagttttga atccaagagt caattatgtt caataatgta atcggccatg 180
 cataaatgtg aaacgaataa agcaacaagc tctaaataac aaatttgatg agaaagctat 240
 tgtgtgatca acgctcttac aaacaaaatt atccacactc acaacggatc gtgagaatac 300
 aacaaagatt ataccataga aaaataataa caaataanaa tttaacatga ttcgacacat 360
 cttgcctata tctacggagc tggttcaaaa atattgtttt atcatataaa taattacaag 420
 aatggaattc a 431

<210> 527
 <211> 330
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 527

tgttgcagac aaaaatctnt ggctagaatg ccgcanaaat ggtctacaaa acatccatcg 60
 gttttcttgg cctgaacact gccgcaacta cctctcccat gttgagtatg gcaggaaccg 120
 ccactccaca tcccgcttgg aaattacacc aatgactgaa gaatcaataa gtgactcatt 180
 gagagatgta gaagacattt ctttttagatt ctccacagaa ggagactcca agcagaatgg 240
 agagatggac actgcagcaa ggcagaagca aattatggaa gcaatcatgt gcagggtctc 300
 ttccactggc aagtctaattg ccagttactt 330

<210> 528
 <211> 450
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 528

tatcaaacca tatacaagga tcaggatccc ttttatatct aaagtaatta tgttcccact 60

gccatcgaac aaattaaagg aatacccaag aaaaaattgt agagcttaag ttgatgttac 120
 tttttcattt gataataaga tttatttggt aaattgaata gatacttatg gttaattctaa 180
 tactctttat ataacacaaa acccatcaat ttgcaggaac ttaatgttcc tgagcatgat 240
 gttgagcagc tattggtgtc actgattttg gataatagaa tccaagggca tattgatcaa 300
 gtgaaccggt tcttagaacg ctctgatagg tcttgccgtt atattttgat ttgttaaatt 360
 aaattcgtca tattcatctt tcttttatat aaaacataat atntactaac atattcacgg 420
 tccaggtcga aaggaatgaa gaagtacact 450

<210> 529
 <211> 416
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 529

aggatagaca aacagcgcta gnccaatcaa ttgtggggct ccaaactcga tgggtggagga 60
 tgcataaatg acaagcaatt catggggctc cggataagat ttgaaggtag aggatagatg 120
 aacagcacta ggcaatcaat tcgtggggct ccagacttga tgggtgatga tgcataaatg 180
 acaagaaatt catggggctt tggataatat ttgaggggtg aggatagacg aacagcgcta 240
 ggcaatcaat tcgtggggct ccagactcga tgggtggagga tgcataaatg acaagcaatt 300
 catgngctc cggataagat ttgttggcag gactgaatgg tccaccggtt tttttccac 360
 cctaaaggcg aacatgtttt atcaaggaan aataaatcat tcatgagagc actata 416

<210> 530
 <211> 221
 <212> DNA
 <213> Glycine max
 <400> 530

gcagaattta gtaatgacct actaacctag aattaatata acttaatgcc attaacctag 60
 ggaattaaaa caaacttaat ggctgagtgt aactgaaatt gtggcaacca aaagtcaccc 120
 ccaacagcca acaagtcagc caccatttgg tctcccaaaa ggctgatgcc taggttgcca 180
 attgggccct tattacaact cgaactaaag cccttttagt t 221

<210> 531
 <211> 440
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 531

 tctgaacatt agctagagcc aggtcaagag gtgagctcgg aatccctgct agaaggtact 60
 gttctgatgc taanactatc tacttgctcc ttttaataat atggaagaac tttagaatct 120
 agttcaatga tcggcgccat tatattgaaa aacatgaatg ctctctttct ttatctctcc 180
 cttgtatata atctctaata actaatttgc caatatgtat cttgccattt attgatcttt 240
 tagatttgca ttaaattggg taccacttgc gtgaaagggc atttgggtct tttgtttttg 300
 ttggtatctt attgctatct aattcgatgc tctcataaag aactagtgc ccaattgcat 360
 catgtctaac tgctattatt tatagccatt gcgaaacatg aatgctctct ttctttatct 420
 ctcccttgta tataatctct 440

<210> 532
 <211> 403
 <212> DNA
 <213> Glycine max

 <400> 532

 tagacggcaa tttcgagcgt ctccatatac tacgggactc aatcagacat ccgagtaaaa 60
 agttattgtc gcttgaattg gcctacaggt tctacattca atatcgagcg tcccgatata 120
 ttacgtcact gaatcggaca tccgagtaaa aagttattgt cgtttgaatt tgctctgagc 180
 ttcaacattc aatttcgagc gtctcgatat attacgggac tcaatcagac atccgagtaa 240
 aaagttattg tcgttggaaat tggctcataa gttcaacatt caatttcgag cgtctcgata 300
 tattacggga ctcaatcaga catcgcgaga aaaagttatt gtcgcttgaa ttggctaaag 360
 gttcaacata taatttcgag cgtctcgata tatttcggga ctc 403

<210> 533
 <211> 448
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 533

tattaaggga tcatttgtcc tttggagtgt tatagctcac tttgaactgc ctaaagtgca 60
 caagaagtga gatcaaaact aaatgcatga gcatgtcttc gccagctct agcttaagcg 120
 ttttaagtttt gatgccaagc tagacatttc cattatgtac tcccttatat tattcttttc 180
 cttatacttc ggagattcat gatcataaat gttattatct ctgccttctt gtttttggca 240
 aagtattggt caatttcttc aagaaatttc tttacatttt caccctcaca aatagagccc 300
 cgaaacgctt atggaataga acacttcatg gtcataatgc acattctatt ggaacgatcc 360
 aatttctcaa ttttggcctc attagagggt ntcgaagtgg atcgttcact tactaaaaat 420
 agactttcaa catcggttat taatcgat 448

<210> 534
 <211> 398
 <212> DNA
 <213> Glycine max
 <400> 534

ctgatgcaac atttggagag gttaatgaaa caacgagatg atgcacttca tgagagggtg 60
 gatcaaatgg agaatataga tcataatgga gaagaaagga ggagaagagg gaataatggt 120
 gttcatagac aaaaccgaat tgatggtatt aaactcaaca ttcctccctt taaaggaaag 180
 aatgatccgg aggcctactt gtatgtgggag atgaaaatag agcatgtttt ctcatgcaac 240
 aactatgagg aggacaaaaa ggtgaagctt gtcgccgagg agttttccga ctatgctctt 300
 gtgtggtgaa acaagctaca aaaggagaga gcaagaaatg aagagccaat ggttgataca 360
 tgggcggaga tgaaaaggat catgaggaag cggtatgt 398

<210> 535
 <211> 405
 <212> DNA
 <213> Glycine max
 <400> 535

tcgtatggtc tgaaacaggt aaaagggcat ggtataagga aattgacagt tattttctaa 60
 aaagaaggtt ttaagaagag tgaaaatgaa gtcactttat atgtgaagtg ataaaaaat 120
 gaagtgcaac tcattgtttc cttatatgtt gatgatttat tttttatata tagggaatca 180
 aattccttaa accaattcaa gaatagtgga accttgaga atctttatga tagatacaat 240

taatgtataa gaaaatgatg gagagctacc atgtcgaaga tcaatacaat aaatataaac 300
aacatttgac agcttctaga aagagaaaac aatcaaaaagg caattgggtgt aagtgggtta 360
caaaacccaaa gtaattctaa tgggttcgca acaactcact agatg 405

<210> 536
<211> 405
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 536

tccgctntca atgtcgagca tctcgatata ttacgggact caatcaaaca tccgagtaaa 60
aatttattgt tgtcagaatt tgcactgagc ttctgttttc aatttcgagt gtctcgatat 120
attacgagac tcaatcggac atctgagtta aaagctattg ctctttgtat ttgctacgag 180
cttccgattt caattacgag cgtctcgata cattatgggt ctcaatcgga catccgacta 240
aaaagttatt gtcgttagaa tttactcata gcctttattt taaattntca acgtgtcgat 300
atattacggg actcaatcgg acatccgagt aaaaagttat tatcatttga atttgctcag 360
agcttctgtt ttcaatttgg agtgggtgctg ataaatgtgg gactc 405

<210> 537
<211> 452
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 537

tgaagagcct cctcaatcaa actgaaaaac ctatattcct caatgaagtt agcacccatc 60
ttttcataaa acttgatggt gagaatattc caatcaagca taaccactt gacccttttg 120
caccctattt ttagggcttg ctttgccacc acagagagca acattctccc aagccccttc 180
gtcttataac actccctcaa gaacaagttc tccatgtaaa accctcgctt ctctagaacg 240
agagagaagt tcggaaaaaa caagacaaac ccaacaatgg aaacaccttt gagggttntt 300
aaattgtagt ttctcgagat aactattact taatgggaat aataaatgaa taattaataa 360
ttatggacta aattataatt gggttaaatt ggaagcagtt ttagagaaaa ctattatttg 420
attggagtag tnggtataag ggtcaatact ca 452

<210> 538
 <211> 390
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 538

ngaggagacg ctgaatcagt tgatgcagat atccatgtcc aactatagga gcacggagtc 60
 ctccaacagg aacctagaga tacaagtggg acaattagcc aaacaaatgg ctgaaagacc 120
 cactggcaac tttagagcca acacagagaa gaatccgaag aaggaatgta gggcgggtgtt 180
 gacacgaagc caaatgagag tgcaaggaga agcagagaaa gctgaaggag accagtctga 240
 ggaaggaagg gcagacaaag aagaagagaa ggaggaagaa gagaagaatg tcttaatctc 300
 tatgaccaan atccagctag cccaagaggc tagaaagaag aaccaccagc cccttctaag 360
 gagcctncat atccttttagt actatcgaag 390

<210> 539
 <211> 334
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 539

tgttcttgtg tgctctcctc cacctagatc cttaattgag ccagtaaaca agtcaaagca 60
 gtttggattg caattgttga agatctgac ttatggaaat tgttgaagag attgtctgag 120
 gcggatgaat ttttcgatga ccgtgatttt cttgattaat ggcaactgtaa ggtctatgtc 180
 ttaatagtga cagctttatt ttattntatt taaaaaaatt gtgcatcaat gtctcattga 240
 cataggactt ttccttgttt gcataaaaact cctttacatt tacgtagaac tctctcgcat 300
 ttgcataaaa tgtagcata tataactatt atgt 334

<210> 540
 <211> 321
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 540

caagcttaat ggatggttgc cttctgttnt acggttcttt gaagggattt cttattatcc 60
tagagaaata ttggtaggtc aatcccgtcc ttgatgattt gcggttggtt tttcctctc 120
gtttcaggaa aaactaaact atntatatgg gtctttttct tttccctctt ttattgcatg 180
gtttgttaat ggatgtacac caaagatagg gaaacaagtc taaagagagt cattgaaaag 240
aaatccctac ttccacggat gatctatctt tcgattaaaa gcgcttcagc atccatcaag 300
gagcatgttg aggtcaatgg t 321

<210> 541
<211> 213
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 541

attcattatg cgaataatg tgctgtaacc cattactaac caattcacat tattaagtag 60
tcgtctggta atcatgacac ttgttggtcc aacaaaaatc atttactggt gcaacatata 120
tgattgtcat aattgacaac acataatgac atgcatgcgt attanagttt gagcgcgaca 180
cacattgact gacttgacta cacattctga gtg 213

<210> 542
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 542

caagctttga gccanaatcc tgactcacca tanaccttga cccagggtga gaatgccaat 60
ccctaccctc ggaagcaaaa aaagaataga ggggaaattt ccaatcaaag aaaaagagaa 120
tgaaaatttc caatgaaagc aaaaaagaaa tgaaggaaaa ttcctcaatc aaagagtggg 180
agaaagcaaa aaaaaggaaa agaaggaaaa ttccccaatc aaagagtggg agaaagcaaa 240
aagaacagaa aggaatattc ccaatcaaag aatgggagac agtataaaaa ggaagaagaa 300
gaaggaaaga atgctcctga tcaaggatcg aaagaaacca gaagaaatgt gcagagaggt 360
ctttggacca tacaatatct gaacagtaca gaat 394

<210> 543

<211> 500
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 543

agcttcttga accgagtagt accgtgctat gctactgtgc gcaacatcgg accacttgct 60
 tgtgcttgan cgactacaca ttgtcaagat gttgcctgag caagatgaaa gccttggaag 120
 aacgaagtgt gcctctgtcg ttgtggatga tatcttcaga gatacctgag tgaactgtgt 180
 ctgaggatat cagacacctt tggactttca aggagttgag tcttatacta tgnatatgca 240
 tatacgtgag aatgaagaca agcatgagtg accatggctg agagtgtgag aacagaccgg 300
 ttactgcatt gtgatcatgt gaaagcatga ctcatgagtg ttatgcatac aatacaccac 360
 atcatagagg catatctgat atgaaacaca tgactatgca tgatgctgct acggtgatgc 420
 ttcatgccaa acacattcgt atatctctcg gtgaagcatg acacagcgtg gtcacgcaca 480
 cagagctcac tgtgagaggg 500

<210> 544
 <211> 375
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 544

atgaagtaac cagctcgctt gagcgagcaa gttactttcg gatgaagcaa gcagctcgcc 60
 tgggcgagct actgtgcaac ctctaccctt catttccat aaataggcat gaggggggct 120
 gaaggaacgg tccaacattt gaaatcaaga ggattagaga gaaatttgcg agaagatgga 180
 gaaaaaaaga agaaagataa aggttgagac gctttcgtaa cgtttctgtg atcgattccg 240
 agatcatttt tcatcgttct tcgaacggat agtttctatt attgaagcta tgaattcatt 300
 ctatgcaccc ttaggggacc atacttgctt tacatatctt catcttcatt cttctaccat 360
 tagngatctt tcttt 375

<210> 545
 <211> 276
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 545

acaacatcct anatagatta tattatattg tagcananat taanaatann tttaaattta 60
agagagctng ataataataa aataaataag aaaaagaaaa aatagaaatc ttactaatac 120
taacagtctc tcaataaata ttctacaaaa attcttgaag atcttcctaa acatttgga 180
agtcttgcaa aaacacaatt ttntntaaaa ataaagtaca nnatttgga nacaagatca 240
ctaagaattt ggaacttctt aagtccaaag atcgac 276

<210> 546
<211> 428
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 546

atgcaagctt gtatattaga caagtcataa gtcacatgt tcatanaaca aatcatttgt 60
ctaagtcact ggcactctaga agtcctaatt ctctcgtaat ggtgtagaac gaatcttgt 120
gtagtgggtc tgtgaagata tttgcgagtt ganttttgggt atctacaaat tctagaacac 180
agtcaccttt taggacatga tctctaagaa gatgatgcct aatttctatg tgctcggttc 240
tagagtggaa aactagggtc ttagatatat ttatggcgct tgtgtgtca cattntatgg 300
ngatgtggtc taatacaatc ctataatcag atagttgtta tttcatccac agaactctgtg 360
cacaacaact accagcagag atgtattctg cttegggtgt ggataatgct acaataattt 420
tgcttctt 428

<210> 547
<211> 298
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 547

accagtcac cctcattcag aagtagctct nttcttcctc tattgcctnt agttgaatac 60
acctttgttt ggttctctta ttggttctta accctctcat gcaacttctt taaaaactct 120
gacctagatt ccccttcttt atgtataaaa gaagtgacta gtgtgaggtg aatgaggtct 180
aacggtgtta ggggatngaa cccatagaca acctcaaaag gggactgctt ggtggttcta 240

tgaaccaccc tgtttaggc aaattctaca tgaggaagat actcatccca agacttat 298

<210> 548
 <211> 376
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 548

ttcaacaagt gtatgaaatg catgtacttc tttatgatga gaaaccactc ttcgtcgctg 60
 acatgttgaa cactntagct ggaaaacact tctttgtgag acagagcaag tctatgcaac 120
 aaagtctctc ttttgatggg gattgaggaa tattagagct tggcttcatt tattcttcat 180
 aagacttggc agatcctact cgaatgtctc tacaaaatag atgttagaca caggattaaa 240
 tgaagtctta aatgtcaact ttaatatcga atcagatcat gattccatct tgcaatcgctg 300
 cgaaatatca gacgtagact ctgcgaaaca tgatntgata acgcatatga tgcaatcctc 360
 ctaaagatgg acccat 376

<210> 549
 <211> 273
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 549

ctgatcaaca catgcacagt ggccaaggat gcatgggaga atcctgaaac cactcatgaa 60
 ggaacctcca aagtgaagat gtccagaatg caactattgg ctacaaaatt cgaanactctg 120
 aagatgaagg aggaagagtg tattcatgac ttccacatga acattcttga aattgccaat 180
 gcttgactg ccttgtgaga aaggatgaca gactgaaagc tggtgagaaa gatcctcaga 240
 tctttgccta agagaattga catganagtc act 273

<210> 550
 <211> 339
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 550

1861 1862 1863 1864 1865 1866 1867 1868 1869 1870 1871 1872 1873 1874 1875 1876 1877 1878 1879 1880 1881 1882 1883 1884 1885 1886 1887 1888 1889 1890 1891 1892 1893 1894 1895 1896 1897 1898 1899 1900 1901 1902 1903 1904 1905 1906 1907 1908 1909 1910 1911 1912 1913 1914 1915 1916 1917 1918 1919 1920 1921 1922 1923 1924 1925 1926 1927 1928 1929 1930 1931 1932 1933 1934 1935 1936 1937 1938 1939 1940 1941 1942 1943 1944 1945 1946 1947 1948 1949 1950 1951 1952 1953 1954 1955 1956 1957 1958 1959 1960 1961 1962 1963 1964 1965 1966 1967 1968 1969 1970 1971 1972 1973 1974 1975 1976 1977 1978 1979 1980 1981 1982 1983 1984 1985 1986 1987 1988 1989 1990 1991 1992 1993 1994 1995 1996 1997 1998 1999 2000 2001 2002 2003 2004 2005 2006 2007 2008 2009 2010 2011 2012 2013 2014 2015 2016 2017 2018 2019 2020 2021 2022 2023 2024 2025 2026 2027 2028 2029 2030 2031 2032 2033 2034 2035 2036 2037 2038 2039 2040 2041 2042 2043 2044 2045 2046 2047 2048 2049 2050 2051 2052 2053 2054 2055 2056 2057 2058 2059 2060 2061 2062 2063 2064 2065 2066 2067 2068 2069 2070 2071 2072 2073 2074 2075 2076 2077 2078 2079 2080 2081 2082 2083 2084 2085 2086 2087 2088 2089 2090 2091 2092 2093 2094 2095 2096 2097 2098 2099 2100 2101 2102 2103 2104 2105 2106 2107 2108 2109 2110 2111 2112 2113 2114 2115 2116 2117 2118 2119 2120 2121 2122 2123 2124 2125 2126 2127 2128 2129 2130 2131 2132 2133 2134 2135 2136 2137 2138 2139 2140 2141 2142 2143 2144 2145 2146 2147 2148 2149 2150 2151 2152 2153 2154 2155 2156 2157 2158 2159 2160 2161 2162 2163 2164 2165 2166 2167 2168 2169 2170 2171 2172 2173 2174 2175 2176 2177 2178 2179 2180 2181 2182 2183 2184 2185 2186 2187 2188 2189 2190 2191 2192 2193 2194 2195 2196 2197 2198 2199 2200 2201 2202 2203 2204 2205 2206 2207 2208 2209 2210 2211 2212 2213 2214 2215 2216 2217 2218 2219 2220 2221 2222 2223 2224 2225 2226 2227 2228 2229 2230 2231 2232 2233 2234 2235 2236 2237 2238 2239 2240 2241 2242 2243 2244 2245 2246 2247 2248 2249 2250 2251 2252 2253 2254 2255 2256 2257 2258 2259 2260 2261 2262 2263 2264 2265 2266 2267 2268 2269 2270 2271 2272 2273 2274 2275 2276 2277 2278 2279 2280 2281 2282 2283 2284 2285 2286 2287 2288 2289 2290 2291 2292 2293 2294 2295 2296 2297 2298 2299 2300 2301 2302 2303 2304 2305 2306 2307 2308 2309 2310 2311 2312 2313 2314 2315 2316 2317 2318 2319 2320 2321 2322 2323 2324 2325 2326 2327 2328 2329 2330 2331 2332 2333 2334 2335 2336 2337 2338 2339 2340 2341 2342 2343 2344 2345 2346 2347 2348 2349 2350 2351 2352 2353 2354 2355 2356 2357 2358 2359 2360 2361 2362 2363 2364 2365 2366 2367 2368 2369 2370 2371 2372 2373 2374 2375 2376 2377 2378 2379 2380 2381 2382 2383 2384 2385 2386 2387 2388 2389 2390 2391 2392 2393 2394 2395 2396 2397 2398 2399 2400 2401 2402 2403 2404 2405 2406 2407 2408 2409 2410 2411 2412 2413 2414 2415 2416 2417 2418 2419 2420 2421 2422 2423 2424 2425 2426 2427 2428 2429 2430 2431 2432 2433 2434 2435 2436 2437 2438 2439 2440 2441 2442 2443 2444 2445 2446 2447 2448 2449 2450 2451 2452 2453 2454 2455 2456 2457 2458 2459 2460 2461 2462 2463 2464 2465 2466 2467 2468 2469 2470 2471 2472 2473 2474 2475 2476 2477 2478 2479 2480 2481 2482 2483 2484 2485 2486 2487 2488 2489 2490 2491 2492 2493 2494 2495 2496 2497 2498 2499 2500 2501 2502 2503 2504 2505 2506 2507 2508 2509 2510 2511 2512 2513 2514 2515 2516 2517 2518 2519 2520 2521 2522 2523 2524 2525 2526 2527 2528 2529 2530 2531 2532 2533 2534 2535 2536 2537 2538 2539 2540 2541 2542 2543 2544 2545 2546 2547 2548 2549 2550 2551 2552 2553 2554 2555 2556 2557 2558 2559 2560 2561 2562 2563 2564 2565 2566 2567 2568 2569 2570 2571 2572 2573 2574 2575 2576 2577 2578 2579 2580 2581 2582 2583 2584 2585 2586 2587 2588 2589 2590 2591 2592 2593 2594 2595 2596 2597 2598 2599 2600 2601 2602 2603 2604 2605 2606 2607 2608 2609 2610 2611 2612 2613 2614 2615 2616 2617 2618 2619 2620 2621 2622 2623 2624 2625 2626 2627 2628 2629 2630 2631 2632 2633 2634 2635 2636 2637 2638 2639 2640 2641 2642 2643 2644 2645 2646 2647 2648 2649 2650 2651 2652 2653 2654 2655 2656 2657 2658 2659 2660 2661 2662 2663 2664 2665 2666 2667 2668 2669 2670 2671 2672 2673 2674 2675 2676 2677 2678 2

<400> 551

<210>	552
<211>	286
<212>	DNA
<213>	Glycine max

ggagatgaag atgattgana agtactgtat ggatttgttt tntcatggcg aatatgtgaa	60
tgtatgtata .catgattntg atgatgtcaa aagaagaatc aaacaaggct catttgattc	120
aagattaata caagattggt tcaacaaata aagccttgat tcaagatttc ttcaagatca	180
agccttgccct canaatgaaa ggtttcaagt catccaaggc acatgtaatc gattaccatt	240
acatgtaatt gattaccaag gcacatgtaa tcattaccaa tacatg	286

229

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 553

agcttggttcg gntgatttct ttgataatgt tttgctcttt tgcttagaag gatttcttat 60
tntcctttga ctttataatg gttctcttgt accgactaaa tttcctcatt tatgagggaa 120
gaatgcgtct gttttatgta ctggtagat gagttcgtgc caagcttgaa tgaattctta 180
agaggggtgtg ttttcaaagn ttatacttta ctctgtatcc tcttggaat ttcaggggtc 240
aagtgccatc ttcagcgtg atctttttgg tgactcaaga gancacaaca tcgatcttac 300
tgctgngac cttatcaatc gattatcttt ccaggtattc tttacctttt actcgaattg 360
ggaaattca 369

<210> 554
<211> 526
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 554

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agacnnaagc nanatcctca gatggctcnt gtaggactag acttatacca acctacacca 120
tgggcacaac atgttgcaat cgaaaactga gtccgcacac ccctaactta agactaagtg 180
gcagtgatgc ttcatgcaag tgctggggca acagtacatt taccaatgct aaagtgacgt 240
aactaggcac acaaatggat gatcataccg agagcataca aacattaaga actgaattaa 300
gcattgaaca caggaaacac agtcaactag atgtacaagt aatgacatta gactatctac 360
agaaatcccc agcaaggggtg ttcagccagc cattacagaa aagctctaac agtgatgaga 420
ttacaaaacc taggcctntt tgcgaaagat gctccacttg ctgcctctag agcggtatct 480
cgagataaga gtagggcgcg ctcttgaatc attgcaaagc atctcn 526

<210> 555
<211> 320
<212> DNA
<213> Glycine max

<400> 555

agcttttact cactgttttc atcagatata atttcccttc acgagataat cgccccctttt 60
 cagattttctt ataatgtcgc aagaatcagc agctccgaat tctgtatttc gattctactc 120
 ttgaagtagt attgacacat gccctttcgt gacaatttta ctgctatcgc cattcttatg 180
 ctcccataat atgaggctat actatgctta tcttaactcc gaacagtgtt gcgttatatt 240
 tatacaaata tataaccattt atactcacag tacgtatgtg aggggttaat tcttctaact 300
 cataagcacc attcgaatag 320

<210> 556
 <211> 330
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 556

gctctagccc tcgaattgtc aacataactt cgctagagaa actctagcga gtacatcgtg 60
 catatggaga acatctngta nagccccctgc acatagacca aataaccatc accaaactat 120
 ataattatgg tggaagtgtg gaacaacata aagatggcaa gttattaaaa tccataaagc 180
 cttatatctt agcagaaaaa ctcagaaatt agttaagcaa acatgttcca catatatttt 240
 gattctatgg tttattaaat aacagaagtg gagttnctct tcatgcacag tacacacttc 300
 aattatgaga cagatgttaa gtaatttaag 330

<210> 557
 <211> 303
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 557

agctntgttc cctttgcttg gatgtttgaa attctccaaa ttagtgactt aaaaaatgtt 60
 catacggcgg gttttggagg ttcacgtgat ataccctctt cttattccca aatgagagga 120
 ggccttactt aaaaccttcc cagcttcctt tccattgcta tctcatccat ccaaacatat 180
 ctagctcaat gagaagggat ccaggctttc attaactatc tagctgttat acaattgtta 240
 tatgtttcaa tgttttttgt gtatttcttt ccattntctt gcccncaact ccacatgttt 300
 ctc 303

<210> 558
 <211> 213
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 558

gcttattatt attccaataa cntgtgtgc ggaacaataa aaataacact cataaacata 60
 ttaaaaagca tttaacaatg aggaanaaaa tgtcatatac caaacaagaa gaatagccac 120
 aagagaataa caataaaata gtatttattt tctaaatcta cctncttatt acctaattag 180
 ctcaatctcg caaaattgaa aatgcacaat tga 213

<210> 559
 <211> 311
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 559

tgcgctaagc cactgctgtc ttgcttagtg ggatggtctc actaatcgca tggttcaggc 60
 ttagcgagac aacactntnt gaaccttcat aattntctcc tttttacttg aanatgaagt 120
 gaaatttaca ttaaattgaa taggaaggct tctagtgagc acaaatgata actaaactag 180
 aaatatttac aatcctacca aaaaataacc ataaattggg agaattatnt acattntgga 240
 cactnttcta tacaaaaatt agtcgtaaaa gacgactaac acatagtcac atatgttggt 300
 atgtaagtag t 311

<210> 560
 <211> 375
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 560

ttcttaataa actnttaatt ggggtgcaata tttatgatga taaattattt cttttgaatg 60
 atatattatt tgtataactt aatanaaaat ttatataatt taagaaataa attgtataat 120
 taacataaac atatttgata aataaatata tcataataac tctcaatatt tgtcttacga 180

gaaatataaa catacattca ctttctcttt catctcattg tgcaaacatc tctctattta 240
 tttttcatta gactacttat actctattac ttatttaata ttgagaatta atgtgcgaat 300
 aactcatgag aattctatat aaatatactt tatactatga aaatatattt caaatactta 360
 ccggcttgga aatga 375

<210> 561
 <211> 335
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 561

tgtatggaga tctgagcact tcttctctct atgtcttcac aatcatctct taggcggaag 60
 ttgcagaatc atcatgatga agatactata ttengcacgt tatcgttgat attttggacc 120
 cttaccttga ttgcattgct taaatatgta ttcatactat tgtgtgctga tgacaacggt 180
 gaacgtatgc tgctgataac tttattctct tagataattg ttcttataca taggttacga 240
 tctgtgtgag tgtgagctga gtatgatact tgtgtcatca ggtggaacat ttgcctttat 300
 tcgctgctgt ggatgcatgc cacgcttaaa ttact 335

<210> 562
 <211> 298
 <212> DNA
 <213> Glycine max
 <400> 562

gatctaagaa taggaaaact taattatcct acttggatga atatgaagct tgaggaacat 60
 ggatagaata agaatgaagg aagaaccggt gctattgact gtttgtccta catggacaaa 120
 tttatcgctt acttaactat gtcaacacte aaccaatatt gattcttctc attgcccacc 180
 accctaccag tcaagaacac ccaatcatcc acaagggcca ccctaaatc agccgcacag 240
 ccctgctgtc ggacatacga tatcaaacac cactcttaac acataccatt acactaac 298

<210> 563
 <211> 270
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations

<400> 563

cttcattcaa catgtcatat gaacaggana gantatacaa attcattaac aatcaagtca 60
cactaaacat tacaggacaa cataagccaa cctaaaatcc tagaatgcaa acctaaaaac 120
cagtctctga attgagcaga cctaaaccct aaacatctaa cttccaactc tggaagccca 180
agaacaaaact tcccaaagat caaatcccaa acccaacctc agaaccacaga aacgtatatc 240
tagctacatc atagacaaac agatgacagc 270

<210> 564

<211> 339

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 564

tcaagcttca gactttcgat tccgagtaag ttcctattct tgagattgat tttattatac 60
ganagtaaag gtgaggggtca ccaccgatat ttatgtttga attggataaa gtcattaact 120
gggtgtattgg actaaaatac tattaatata ttttgacata acaagttgta cacatgctac 180
aagattatgg cagaaattat cctgatagat tttttaaaga tttatgatag atttatagtt 240
cattatattt aatggattat atgtgagaat caaaacattc gaagtattcc tatgaatctg 300
cacgatctga aagagtttta ttaattctct ttttaacaaa 339

<210> 565

<211> 515

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 565

aggggnccat gacccatag atangcttag nattnttcaa ancattttgc aagaaantct 60
gctangacta tatatatgca tttttgtgta tgaaaacaag ttatcagatg aagctaggag 120
taggtggagg ggggtttaac tagtaattac cgctcttatt cacaattaat tagccttata 180
tattttttcg ggggacggtt ggtgctggtt cgaatcgaat gaaatctctc ttctgtagtt 240
tgccaaaaat agtttcttga gatcactttt atatttttta tttgtaattt cattcttggt 300
gaaccgtgaa ctacatcacc atcttctatg aataaattac ctctaattta tgggcttatt 360

ttgaaacagg ttnttatcca ttgcttcttt cnaccagcgc gttggtcctt ccacaagaat 120
 gttaaacagg aacgngata gagggctctcc ttgtcttaag cctctntggn gaataaaactc 180
 aggtgtagga cttccgttaa ccaatatgga tatggaagca gacttttatgc agccttcaat 240
 ccaagttacc cacctgtcac anaaacccat cctcttcaac aaatatagta caaactccca 300
 agatactgaa tcatangcca ttntataatc tactttgata 340

<210> 569
 <211> 240
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 569

caattggaag tgatcatgga ggtgaatntc ataatgagtc ttttgaacac ttttgtgaag 60
 aacatggaat tcaccacaat tcttattccc caagaacacc tcaacagaat ggtgttgagg 120
 aaaggaaaaa tatatcccta taagaagggtg caagaaccct tctaaatgaa acaagggttac 180
 cgaagtactt ttgggcagat gttgtacata ctatntatta caccttgaac agagtactta 240

<210> 570
 <211> 351
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 570

tgcacaagac tctttaatat tgaagagtat ccttgtggaa ccttcttccg acgaagacac 60
 tgacaaaaac ttatcttctc cttcttggac aaggatgtg caggctgggg caagtaaatt 120
 ttcttcccat cagaccttgg atgcaattgt gatcgtatac ccatatcagc tagatcttga 180
 tgggtattca agccatcctt cgtcttgcct tgaatgttaa ggagcgtnc aatcacattg 240
 tcacaaacat ntttcttcac atgcataaca tcaatacaat gtctaacgtc aagatcacac 300
 cagtacagaa gatcaaagaa natagacctc ttcttcatat gcaactctga c 351

<210> 571
 <211> 421
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 571

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acattcaaaa ctggaaactt agattcctag ccatgagtca tccttttggc actgtagttt 120
agcttctaca aactaccac acactcaca tgtgcacaat ttgtttcgca agctaaattc 180
cacaaaatca tccgcaaagc ccattgagggc atttcaccga acacttggtg ggcattatgtt 240
taagcatgaa aatcaaggga atgagggcaa tgtggcttgc cccattatct cagaatgcac 300
cctatgccta aggccatacn ctacaacccc acaattcaac aaaaacaagc aaattcaagg 360
atacatccct tcacgtttga gcaaataat gcaacttaga gcacaaaat atatcaatgg 420
a 421

<210> 572
<211> 232
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 572

atatcagtat aactctgata tggcttacga cggactcag ctgcagaaca tggttaaaaa 60
ggatagcgag tccttttaaag agtacgctca gctgtggagg gacctggcag cgcacgtagc 120
ccctcccatg gtcgaaaggg aaatgattac catgatggta gacaccttgc cagtgggtta 180
ctatgagaaa ttagtanget acatgccctt cagcttcgca gacttggtat tc 232

<210> 573
<211> 317
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 573

atctagaggt atacctagga tctctatcag acactatgct agatggcaca ccatgtaata 60
tgacaatctc actaatatac agacagggtca actnttccaa ggaagatatg atattaatgg 120
gaataaagtg agcagacttg gtcagcctgt caacaataac ccacatagaa tcaaaacctc 180
ttgggggttct aggtagtctt acgacaaaat ccatagaaat atatgtccat tgtcactggg 240
tatcttcaag ggggtgtaact atcctgaagg gctctgatat cttatactta tgacagacta 300

aacatgcata cacaaac

317

<210> 574
<211> 180
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 574

agcttgaaaa ataaatattt aataaaaata tatttattta ataaaaatat taatttttgt 60
ccaaaaaaat tattaaacta aaataggtgt taatttaaaa ntgggctttc tgccttaa at 120
aagctggacc ggccctggaa tgaaataatg ggaaataaag aaatttggtg aaaagtattc 180

<210> 575
<211> 380
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 575

gctgccactc aagagacatc tcacctatat ctntatttaa aactatcatg tacatctgtc 60
cattattcan aaataacacc atctaagcaa acttaactga gtagagacta gtactctcct 120
tcttccatac caatatgtcc tctcaatca gaatcaaatt aatacttcta aagtcatgac 180
cttntatcta agtaaattat tatatntatt tctcctaag atatagagat ttctncattc 240
ccatcaccac aatcaatntc cctccctccg gtcacctaaa ctttgacacc ctggctttat 300
ggctactaca anggtgtata atctagtata cttcctatat gtgaaaataa aaccacttca 360
cccgaacgtc ttgagaaaat 380

<210> 576
<211> 147
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 576

tcaagcttct tctggacctt gaacaagcaa tcaactttct ctgtcacaac catgctatgt 60
gctcgcgact ggctcccttc ttcccttcgc aacttgagct cactattgct accccataga 120

gctncgagaa atttggtccg gccatac

147

<210> 577

<211> 324

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 577

caccggtcac gtgtctacta tcattgttat aatctctntc tctggtttgg ggggtgtact 60

tgagttgcc aacctctcca tctttgngcg tggtctttga aagatccgtc cncctattg 120

cacacgttct gtagttgcat cctatccgaa gacattatac tgacactgcc taacgaaggc 180

caccactagg tccttccaag aatggactcg ggaagggtcc aagttagtgt accaggtaac 240

agctaccag taagactttc tcaggagaaa tgtatcagca gtttctcatc ttttgcgat 300

gcacgcatct tccgacagta catc 324

<210> 578

<211> 215

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 578

gatccanaat cctgactcac catanactct tgaccaggg tgagaatgtc aatccttatac 60

ctcggaagca aaaaagaaaa gaaggaaatt tccaatcaaa gagaaagcag agaaaaaaaa 120

aaaagagaag ataggatatt cccaatcaaa gagtgggaga aagaaaataa aaggagacga 180

aagaattttc ccaatcaaag aatgcgagaa agtat 215

<210> 579

<211> 270

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 579

ccaaaactca ttccgtagat ccctcttgta agactaagtt tcaatcttgc ttcaatcaag 60

ttctaaggca acagtacatt tcccaatgct aaagtcacct aactatgcac acaaattggat 120

gattagacca aaagcataca aacattaagc attgaacaca aaaaacataa tgaattagat 180

attaagtatt tacatcagtt gctcattaga aatccncaac taggggtgnt agccagccat 240
 tacagaagag accctaacaa taataagctt 270

<210> 580
 <211> 440
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 580

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 aataaaaaga ataataaatc cacagcaaca accttatctc tgtagccgctc aacaccaatg 120
 ggcgaggtct gtataaccat tctctnttcc ttttcttttc ttcaattacc atcaatgtat 180
 cattccgggt tctgattttt ttttgtgtta taaatacgaa gagaaaaaac tagaggaaaa 240
 caaagtggaa gagaaaaaag cacaggaaga agaaaagaaa gaagaagaga caaacagag 300
 gaatcaaaag atggcaagaa atccaacgag gaatctgctc caccagaaat cgtgcaaggc 360
 acaaccttca caatgcatga acactttaag cagatttctt gacatctttt ttaagntaag 420
 gactaaattt gcacacttat 440

<210> 581
 <211> 368
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 581

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 gaactttgag ttgtgtctca caagactctc attcatcaga gttacaacaa gtgttacata 120
 tgcttctata tatagactat gtagcttctc tgagaagctc tcttgagaaa acttccttga 180
 gaagcttctc tgagaaaact tccttgagaa gctagagctt agctacacac acccctctca 240
 taactaagct cacctccttg agaagcttcc ttaagaagat tcctagagaa gctagagctt 300
 aactacacat acctttctaa tagctaagct caccttcttg agatgagacg ctagaactta 360
 gctacaca 368

<210> 582
 <211> 307
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 582

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 aggaaagatg cgcggagtcg ccaccaacgt ttatttgtgg aaaacgtctg ataaaaccga 120
 aggaaactgg tcaaaaggaa aattctaagt tcgggagttg tatttacgct tgaggaaggt 180
 attagcacct cacacgtttg tcccatagga caacagtcta ttttttagaa ttgcggaatt 240
 gtgttatctt aacctttagt tctttttatc ttttgaggtc aacanaagcg gggcttttgc 300
 tcctaca 307

<210> 583
 <211> 361
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 583

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 ccagaatcat aagagaatat gaagatggag gaagccgcag gcagtggcaa cagtgcggt 120
 ggcaacaaat ggtggtggng agtggccagc gctgctcana tgggaatgng aattcgtacc 180
 ttcgccaaag ggcacggagg cgattcgcgc ctcatgcctt tcanagcctt cgttgtagct 240
 tccctcttcg tcagcagcgc cgccttcgcc ttcgttctac tctccaagc taacggcatc 300
 cacagggtac tactgctctc atcttcattc ccatttctgg tttctaactc tctntatgtc 360
 t 361

<210> 584
 <211> 201
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 584

catcatcatt gtaactatag ctatccatta taaaatagct tagacaatct aggtctgctg 60

cttctaaacc ttntatttat ccttgaagca gctcaatggt ttttatttcg cgtcaactag 120
 tgaagagtga aactgaagc tgttctgcag actcattgat tttccttcat tgcgacagaa 180
 cctaaccata ccctgccctg a 201

<210> 585
 <211> 198
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 585

acacaccctt ttatactaaa tgcancctccc ttttctattt tgtttgaaat tctttntccg 60
 taacgttacg aaactttacg aatctcgtaa cgatacttat tntccttccg cacggttacg 120
 aatccttacg gattatgtat ttactctnnt ttggctttca aagaagttac ggacactcac 180
 ggattgcgca aaaacacc 198

<210> 586
 <211> 399
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 586

agcttggttn tgggtcaatag caccctacct gacttcnca aggtctactg atcctcgcaa 60
 catatctcca ggtaccactc tgtgggtcaac aaataatagt aggaagattg actcttccat 120
 gctttctcac atcaagetta ttggattatg gcgcaccctg catatgtggt actaggtggc 180
 aatcgggcga tggcacaaat caactatcac atttccacaa gccaggcata agcacaccat 240
 ccncagctgc ccacctttaa atttagctca cgtgcatgta cgtagccttc tcctcgttcc 300
 tctcagcacc ggggtcccat caacccctc aagctntcac aatatccaat caattcaatc 360
 ccatttgta tgaaactacc ttaaacaaag aanaacaga 399

<210> 587
 <211> 314
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 587

gcttgaatcg atacacacat actgtaatcg attaccagag cagtatttca gattatattc 60
tcatcagtca canntctttt attggttctt gaatggccat canaggctta tatatatgtg 120
acttgagaca ncgaattgct aagagattnt cagaacanna aggtcttata ctcttananaa 180
gcanaatcgt ttcacacctt taanaattcc ttggccaaaa cacttgatgat tcaataagga 240
attagttgag tgctcaaatt gttcaatcta tctcttcaa gagagaatac ttcttctctt 300
cttctntatt ctga 314

<210> 588
<211> 383
<212> DNA
<213> Glycine max

<400> 588

tgagagcgcg atcttatact gtgagagaac gactagctgc gagtaataat ctttgcata 60
atctctgaat tctagaatga aatgtataaa tgaggacatg atgaatgcta tgattgcaca 120
tacacaaggc ttttgaccaa aaagcttacc ttgaatgata attatatacct tcgcaccctt 180
tatgagctga atgatattgt caaagatttg aaccctgaac ttaaataata acctccagat 240
accttggttag attctaggag agcatatggc tcaaggcaaa ttaccgcaaa tttgcggagt 300
ggaactaatg ggatgcaaga aagaaataaa catcggcaca acaacacata tgttggtgat 360
aacaataaag gagaatgaaa agt 383

<210> 589
<211> 425
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 589

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gtgttgtaaa agccattcaa tttaacatat agttgagaaa ttgatcatca aatgtatgtg 120
catganaaat atatatgttg agagatattg gccttttcat gattagttnt ttaatccctc 180
ccagaagcta ccatggccac ccataatana tgccattcta aatttttgag tttcttaciaa 240
attaagtatg gacaatggac ctaagtgaat ggatttgacc tacgatagtt gatataattct 300

tgatagttaa tatctgaaca ctcaattatt ntataactn catcaataat cattttttct 360
 ctttctatca tatcatctat ctatannntt tttttctttt tgggtcattc tctctggctc 420
 tatct 425

<210> 590
 <211> 318
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 590

agcttgtagg attatggtgt acccatcaca tggtgtacta ggtggtgggc gggcgatgga 60
 gcacaaaaca tttccacat ccacaatgcg cgcataaacc caccatcenc tggtgcccac 120
 ctccaactga gctcacgtac tcccacgtag cccatatact cgtttctctc aacaccgggt 180
 ccccatcaat cctcccaagc ttccacaaca tccaagcaaa acaacattcc aacagcacia 240
 gctatcacag ccaagcataa cagagcagag gcagaaaact ctgctcaaca catcaaccaa 300
 aattacagct tttctcac 318

<210> 591
 <211> 357
 <212> DNA
 <213> Glycine max

<400> 591

catgcatgct tacattccct ttagcattca ttattgaatc atttcagcct ttgctttcgt 60
 gtagcttagg aaaaacgcca tgtattctcc tttctttctt ccaaagccat ttctaactg 120
 ccaagcactt tctccatcac ccacatccac cattagccac cacaaacct cgttgctctc 180
 cgggtgaaacc ccacaccgat aggaaccctt caaccaaagt ggaatcttac aacttggctt 240
 gcgggttttg tagagaacga aaccctaate tgaccttttg tttttatcga gaggattgcy 300
 ttgaattgat gagcaacgac gaataagaat cttcaagtga cgcgacgagg aaccgc 357

<210> 592
 <211> 172
 <212> DNA
 <213> Glycine max

<400> 592

THE

<400> 593

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<210>      594
<211>      495
<212>      DNA
<213>      Glycine max

<223>      unsure at all n locations
<400>      594
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<210> 595

<211> 390
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 595

 agcttaacca agaaaaaggc taacaatggt nttatgcaca attgaaggaa ataaaattca 60
 gaatttagga attcaagtaa caatccttca tacaaccaat atattacctt aaagagattt 120
 ttttttanag ttcttcaagc atcaaccatt caacccaaat ttgtctctct tntttttttt 180
 ttttaattnt gcttatacga atttctgttt tttttttata acaaagagat caaaaggctt 240
 aacttttgca atggttcagc ctaaaaaaa aacatgaaca agaaggtaat ataaatggca 300
 aagaaaataa agaaggatgt tacccaatat ttccagcaaa ggaagtgttg atcctagaat 360
 cggaactctg ataaccaa at gatatgaacc 390

<210> 596
 <211> 318
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 596

 cacaagtgag tttattcgat tattagagtt atctctttat cttaggagag tgattctctn 60
 attcttgggtg atcaagacac tctgctgac naggacttca caccttgtgt gtgccctcct 120
 ggnagagtgt tctttcttct atcatctcac ctgtctttt accacaattc agaaatcacc 180
 ttgccaaata tctgtgacat actccattac aactcaata agtattttga cctaatgaat 240
 tcaaacagac ttactcgtt gaatacctca tggacctgac ttcgtagtga tttatatttg 300
 tcgcacacta ccacgtta 318

<210> 597
 <211> 396
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 597

 aagctctact aaataagcct agacattaaa attttgatag anaaagtcac atgcatgttc 60
 cctgtatata gacaagattt gagttgagat gaactgaaga aagagagggc atatttagta 120

ctctagctag tcaaagtggg atatatatgt ctaataacga acctggtgct ccataatttt 180
 tgcaccccggt gtccgtgtttt gttgaatctc tccctccaac ctaaagttat atagtcagtc 240
 acaaaaagag tagtaacagg gttaagcata tatatgtgtg ttatgtcagt tatctttcac 300
 aaagcagtat atgctatata tatcgctcac agatcaaagc tgacgatagc taaacttact 360
 ccaataggtc tgatgcaaga acataactcat tcatgc 396

<210> 598
 <211> 511
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 598

cggaatgaac catgtagnac ccataactat tacgtaataa gtctacagaa ataccagtca 60
 tgcatacttc caatcaatcc acggggttat ataacatgca aggtgtttgg gccactaaca 120
 tgaaccacat tcataaccca cctgtcaaga ttctgattaa tcaatgctgc tgcaaatctg 180
 caaaactcaa tatattgaag attgaagatt ctcataggaa aattcctaaa tgcatactta 240
 taacaagaaa gtaaataaga caaaaacaat accctccaaa acctgctcgc atgtccataa 300
 catttcttaa tctgatttct ttccagtgtg aaacacgaac atagcttgct attatttcat 360
 tccagtattt cgaatctgcc ctaaaaagct ctgatctgga tgtgaaagca tcaagctnta 420
 tgctttgaag cctatcatgt ggggtttgca aacgtgcatg ccattagtaa catttgctca 480
 tatccgttct tatgcattca gagatgcatg t 511

<210> 599
 <211> 415
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 599

gagggactca tgggcactat gaatgacaaa ttccgtggga taaaggtagt ggtgccatgt 60
 tcccacagcc cgtactaagg catacaactc cttatnataa gttgaatagt taagggtacg 120
 accacttaac ttttactaa aataagcaat tggatggcct tcttgcatac acacagcccc 180
 aatcccaaca tttgaagcat cacactcaat ttcaaaagat ttttgaaagt ttggcaacgc 240

aagtatggng gcattacgta gcttttgctt aagaacattg aaagcttctt cttgtttctc 300
 tccccatttg aaaccaacat tcttcttgag cacttcattg agaggtgctg ccaatgtgct 360
 aaaatccttc acaaatcgtc tatanaaaact tgctaagcca tgaaaactcc tcacc 415

<210> 600
 <211> 451
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 600

tattgtagcc gatgctctgt ctcggcgctca tgcattactt tctatgcttg aaacannaat 60
 gattggctctt gaatgtttga aaagcatgta tgataatgat gaaacttttg gagaaattct 120
 taaaaattgt gaagaatttt cagacaatgg tttctttaga catgaacgct ttcttttcac 180
 agaaaacaaa ttgtgtgtgc ctaaagtgtc tactagaaat ttgcttgatc gtgaagcaca 240
 tgangagggt taatggtgca ttntgtggtc caaaagactc tatagacatt acangaacat 300
 ttnttattgc ctcatatgaa aaaggatgtg cagacactct gtgaacatcg cattgtattg 360
 taaaatgcaa gtctaattgt aagcctcatg gattgatact ccattgcaat accgagtatc 420
 ttgattgtta tcatggattt gtttgggctg c 451

<210> 601
 <211> 322
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 601

tgcgcatact tcttcacgaa cgttcacttg cacaagacat tcttataact atgtaaaatg 60
 aacctatata ctattaatgc accttcgtta cctagattat ttacatgtac tatcaacgtg 120
 tatgtgttac ctacatcaca cacattttct ttgctagact cacatacatg catactctaa 180
 gcactgtggc tatcanaaat tgcatacgtg cacatcnttg gatctctaata acctatacat 240
 acacaaactt cataatgaat cttgactatc tacacaataa ggcgctacat ttcatgctgt 300
 ctttcaagtg ttgtgactac ct 322

<210> 602
 <211> 279
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 602

 gatgatgacc gataacccaa agaatgattc gaagaatgag tcaacaagtt caagtttcat 60
 gagaaganat caagaagatt caagaatcaa gagaagtttg atttcaagat tcatgagaag 120
 atgaattcaa gtttcaagag aagaaatcaa gaagacttca caagggaagt attgaaaaga 180
 ttttcancaa acaaacatag cacagttctg ttttcanaag agttttcttc acaatttcta 240
 gtaccagagt ttactctct ggaatcgata ccagttcct 279

<210> 603
 <211> 272
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 603

 gtctcaagac tggactaata catntgctgt ccaagtttta tggctcttgca ggtgaagatc 60
 ctcataagca tcttatgaag ttccatattg tctgttccac catgaagcct cctgatgtcc 120
 acgaagatca tatctttcta aaggtctctc ctcatctctt ggagggagtg gcanaagaat 180
 ggctgtacta ccttgcttcc aggttcatta ccagctgnga tgaccttaag aggggtgtct 240
 tggggaaatt ctccctaca tctatgacca ct 272

<210> 604
 <211> 512
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 604

 tggggnatga gactcctgat nacgtcttg antttacgga cncatttcaa tactcaagct 60
 tgtgaagtat gtatgttaat gaggaggagt ttaatgtggc attattggac ggatanacgg 120
 ggtgtcttag taggatgggt gtgcccgcct acacactcat taatatttta taaagtgtaa 180
 aatgtatgc ttcaacattg aattctgaga tagagcacga agataatatc taactgatgg 240

cttaaattta ttgataagag ggagtagtct atagaaaagg tgtactttat acgatgaagt 300
gatttttata aatgagataa tggcttagtg eggctaattg tgataataaa gaatttggtg 360
tttgtgaaca ctattgtag gacattgggg atatttttgg ttctaatact gtaaacgatt 420
acatgtaa at tttctattgc ttatatgata aggctgaatg aataggtgta aattaatatt 480
aattataatc ggagttttat gtctgtatct ag 512

<210> 605
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 605

ataatgatgg tccgagntat gttgtggagc ggntacgaac ccggaatggg tttaggcaaa 60
gacaacggcg gcataactag cctgataaat gccaaaggaa accgtgggaa gtatgggtta 120
tgctataagc ccactcaggc aaatataaag aggagcgttg ccggaaggaa gagcgggaagt 180
caaggctcgc ggatgagaca agaaggtgaa ggaaaccac cctgccaaat aagtaagagc 240
tttataagcg cgggtctgga ggacgaaggt caagtcgtca ctatatatga agatgatgct 300
ccgagtacgc tgtatttggg acgaccatgc cttcttgatt tacaactggg aaatcggcga 360
gaggaggaac gccccgactt ttacgcgaag agca 394

<210> 606
<211> 369
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 606

tttctctaca atngcatcac ctctcaatga gctgggtgaag aagaaatgtg gcattacctg 60
nggtgaagaa caagagcaag cctttgcttt gctcatagaa aagcttacta aggcacctgt 120
tctagctctt cctgactggt ctaanacttt tgagctagaa tgtgatgcct ctggagtggtg 180
agttggagct gtattgttac aaggtgggca ccctattgtc ttatttagtg aanaacttca 240
tagtgccacc ctcaactacc ccacctatga taaagagctt tatgccttaa taagagccct 300
ccaaactcgt gaacattacc gttgttcaag gaatatgcat tcatagtgat catcaatcac 360

ttaagtaca

369

<210> 607
<211> 367
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 607

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agtatgatag tcaccgcttt aggagcgagg tacaccagca gcgcttcgaa gccatcaagg 120
gggtggtcggt tctccgggag cgacgcgtcc agctcagga cgacgagtat actgatttcc 180
aggaggaaat agggcgccgg cgggtgggcac cactggttac tcccatggcc aagtttgatc 240
cagaaatagt ccttgagttt tatgccaatg cttggccaac agaggagggc gtgcgtgaca 300
tgagatcctg ggtaggggt cagtggatcc cgttcgatgc cgacgctatc ggccagctcc 360
tgggata 367

<210> 608
<211> 280
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 608

cttatcaagt aaatggatca ttcttaacgt ccaacgcctt ataatgatca cctttcangt 60
aaaaagaatc gcttgattca cgcttaagaa agaactacat aggttcgatt tcctcatcga 120
tgtaggggtac gtatgagcaa aagccccgct attgtcgacc tcataatata aaaagacata 180
atagttaagg taatacatat tccacaattc taaaaaatat gttgttgtcc tttgagacaa 240
acgtgagagg tgctaatacc attctcaaac gtatatacaa 280

<210> 609
<211> 394
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 609

agcttgattt tcaaattatta tgggtgtgcgc ttgttgtaac atgttatgtt tgctactgat 60

ttttaattct ttgacccttt gaatgaccaa attggcttcc gatgtcttca tgagacttgt 120
 agagaatttt atccttttaca ttcaagcact ggtatcatgt tatttggacc attacaacat 180
 aatcaatcct tanagcattg cagtnttggt atattgtgag gacaaactga catctctatc 240
 ttcatgggtca gtttcttcca agatccaagc cttatttgcc catgacttct ccataaaaga 300
 tatatatatc tttctcttag ctntctacaa ccaactgagat catcccaaatt tcacttttgt 360
 agctcaagta gttntcaaatt tattgcacac atat 394

<210> 610
 <211> 525
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 610

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 tcaagtgcgc catgccnctt gatatacttg tgggactcat ggtcactatg aatgacaaat 120
 tccttgggat aaaggtagtg ttgtcatggt ttcaaagccc gtactaaggc atacaaatcc 180
 ttatcataag ttgaatagtt aagggttagga ccacttaact tttcactaaa ataagcaatt 240
 ggatggcctt cttgcaacaa cacaacccca atcccaacat ttgaagcatc acactcaatt 300
 tcaaaagatt tttgaaagtt tggcaacgca agtatggggg cattagttag cttttgctta 360
 agaacattga aagcttcttc ttgtttctct cccatttga aaccaacatt tttcttgagc 420
 acttcattga gaggtgctgc caatgtgcta aaatccttca caaattgtcc ataanaattt 480
 actaagccat gaacacttct cacctttgtc acggacttan gtgag 525

<210> 611
 <211> 386
 <212> DNA
 <213> Glycine max
 <400> 611

atgcaagctt gcagatagat caatgtgagt caacttttat ctttgatcaa attataaatg 60
 tttgaattgt tcttaaaatt ataataaaat caaatatgat aaaataaaaa taaattctat 120
 ttctgaaaaa aaaagtcaat tctacttta cctattgaat aaaattatct taattcagaa 180

ttaatttttt cactactgct aattcaaaca cacacttacc ataaacacgc gcgttgcaact 240
 cgaaaatcaa ttgtctcccg ctccagcaaa atcaaattag taaagcgatt gccacataaa 300
 tttatagtaa caaataacaa tcatcaatgc ctcaaagcta aaccccaaac catctttcttc 360
 acctttttctc tctgcacacg aatatg 386

<210> 612
 <211> 273
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 612

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 tctaattctc ttatagtntt aattgtgtaa tcttgattgt ataaattatt aaatatataa 120
 acatttgctc ttattttatt ataactatata gattgtcttt acattattgt atatcattta 180
 aatattatga ggatatgaaa ttataattta acctttataa aaatagatgt aacgcaacat 240
 agagactgat gctactttga tattccaatt gat 273

<210> 613
 <211> 273
 <212> DNA
 <213> Glycine max

 <400> 613

tatcttgtgc attcaatatc ctgatgaggg tgtttcatat gttctcaaga ctggacaaat 60
 acattggctg cccagtttct atggacttgc aagcgaagat cctcatacac atcttagcga 120
 ggtaaatagt gattgatcca ccatgaagcc ccctaattgt caggaagatc atatcttact 180
 aaagtattta ctcatctctt ggaaggagag agaaaagaat gcgtgttcta cattgctgat 240
 agatccatct ccaactggga tgaccataag aga 273

<210> 614
 <211> 353
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 614

gagatcatcc nctcaacaac attatgggtg atatctcana tggggtaaca tctagacact 60
 ctcttaaaga tntatgcaat aatatggctt ttgtatccat gattgaacac taaaatataa 120
 aagaagccat aatagatgat aattggatca ttgccatgca agaagaatta aaccaatttg 180
 aaagaaataa tgtgtggaaa ttagtagaac aacctggaaa ttatcctatc atatgaacaa 240
 aatgtgtttt tagaaataaa ttatatgaac atgggtataat tattagaaat aaagccaggt 300
 tagtagcaca aggggataat caagaagaac gaatagacta tgaagaaaca tat 353

<210> 615
 <211> 122
 <212> DNA
 <213> Glycine max

<400> 615

tgctattgta acatccactt tattagcaat tgtggaatag attagaaggc cattgatggt 60
 gacgctaata ttgaagaagg aacaattcta ttatagaaac acaaggggaa ccattttttac 120
 ac 122

<210> 616
 <211> 345
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 616

cagtgtttat attgcggttc caatgtggcc agaggggttac ccagaacatg gaacggttca 60
 agcaatgatg gattggcaga ggagaacaat ggatatgatg tacaaggatg atgctggagc 120
 actaaaaggc aagggtaatg aggaagatcc tctcaactat tcgacattct tctgcctagn 180
 taatagggag ctgaacaaag aatgagagta tgtgccccca gaaagaccag atcctcatac 240
 agattatatg agagcacaag tgtcccgacg ctttatgatt tatgttcatg ccaagatgat 300
 gatacgcatg tctgttaatt aagtttacca catcttccaa gatac 345

<210> 617
 <211> 288
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 617

ttctaacaca ggtattttca cggattgtga ctgggatgct cttatactan gaagtctctg 60
cttatatant ttacttttaga ttaaggagga cagatactaa tgctcctttt aacttgaaat 120
atccttcctt ccttcatcag agagcaagaa aaacttacta atggatggaa gcatctatac 180
catgttgcac ntacaatana agcaaaacac acaatctcaa tataacttct attcaatagt 240
tggccaaaga ctttcacaat gagaagagat tagatattaa tataaaga 288

<210> 618

<211> 459

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 618

ngcaagctnt ctaaagtgtt ctttgcccag tctcaggtag agtacttagg acatttggtt 60
tcgcacggn gagtggagcc cttagcatca aaggctactg caattcagca atggccaatt 120
cctcgtacaa cgaaggccgt aaggagcttc ctgggccttg ctggctttta tcgcaggttc 180
attcagaatt atgccattgg tgtggcccca ttagtcaaag ccacgaccaa agaaccctg 240
cattggacat ctgagacaca tgaagccttt gacactttga aacatgcctt gtcaatagct 300
ccggtgttag ctttaccaga cttcaacctt cccttcacag tcgagacaga tgcgtcagga 360
gttggtatgg gtgccattct ttcacagcga ggccaccca tagcattttt cagcaaacct 420
tttagtgcca agtactctga tcataacata catgcgaga 459

<210> 619

<211> 492

<212> DNA

<213> Glycine max

<400> 619

cggaatgata cactttacta tctataatct cagctctcag gagctgagct agttattaaa 60
ggggtgtgtg tagcttatct ctagattctc aagaaagttc tctcacatat tgttctcaat 120
ataacttctc aatgaacctc cctagtctat gaatagaagc atgtgtcaca cttgttgttg 180
ctatgatgaa tgatactttt atgagacaca ctacacagtt ccacttgtct tcctctacta 240
taacttaaac tgaatctacc cctgctcttt ttcttttctt acatttaagc atactctata 300

tgcttcttat ccaagacatc actcttgctg gtattcctct tctttcatga gctacatacc 360
 tatcggctctg tccaggccat atctatcttt cttaacactt tgtgcttcta tagttccaac 420
 atttcccttc tggtctctct tactctcttc atactgtctg ctgtggcctt taatgttcct 480
 ttctccttac cg 492

<210> 620
 <211> 416
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 620

ctaagaagaa agtagaaaac atatcacctt tctatttaac tatcttttgt attgtatgct 60
 tggcaaactc gaatgggtact ttctctctgt tgggggtggaa cttatggatt tccatttccc 120
 aaaaggggtga ctcttctgtc acttccatct tcatatgatt ttggtatgtg gcttcagttt 180
 tagccattat gtgngtctg tggcgcgcca attgtttgac ttcaatgata aattgtgttt 240
 cttactcaac atcaattcat tggaaattaa tgctttcctt aactcagttc tatttgcaag 300
 aactagagaa tactatacac gtgtttctgc atgaacacca gctcatgcag ctacttcatt 360
 aacgcatata taattaaagt tggctctcag aatntctcca tatcaccaac tatgat 416

<210> 621
 <211> 328
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 621

atagcccag gactagagta cttgcataag ggatgcaaca ctggaanttt acattntgac 60
 atanagccac aacattcttt nggatgagaa gttctgctc aagatatctg attntgggct 120
 agcaaagcct tgtcctagaa atgaaagtat tatttccagg tctgatgcca gaggaacatt 180
 atgggtatgta gctccagaaa atttggcaga atttcacaca natctgatgt aatcttcac 240
 tttanattta aaccacctaa accttaatgg gtaaaattaa ttctattatg cattanatgc 300
 atcttatctt tgacttgaac tctacaat 328

<210> 622
 <211> 439
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 622

ttatagacga atgtttcact tttcttccac ttctaccgag agacataaaa ttttcaacgt 60
 ctttcatagc atcaaaactga aaatctctaa ctgcccttcc aataactaatc gctgcctcaa 120
 caattccttt ttctcccatt agcttgctcc tgttttagtga cacaatgcga atgacaatgc 180
 agtttccaaa gtacgttgaa ggaatcgaaa attcgggacg gttacggcaa tctcccagaa 240
 acttcaagct gtaaatttca tcattattat tattatttgc aatagtata gcattgacct 300
 cttcttctga ttgaaccttg caaaccacaca tcanagcact tgtcataaca aaggttgata 360
 tgtgcaatga ctctaattct aaccataag tacttctaca ttcaatcgac acccatttct 420
 ttagcttcgc aacatggcc 439

<210> 623
 <211> 385
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 623

taaacctctt aatttagtca atgcaatttc ttttttatcc cttgttctaa ttacatgcat 60
 aatgcatctg gttntcctag tttttcaatt gttaaaatta ggtccatgat ctaatgtatt 120
 ttttgacatg atagtggctt gtctgacttg ccacatgata canagcaatg gcatgagttc 180
 actggttata gccactgttt tgtctgtcat tgtaagaatg nngaaattgt gaatatgaat 240
 aggatttgaa caagaatctg ttcttgact annaaatatt atttttatga cacattntan 300
 aacggtctaa tcgcttanaa tgtcagcttg tcaatttata ataatanag tatataatga 360
 caatttctaa atgtcttaca tggta 385

<210> 624
 <211> 335
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations

<400> 624

gtgtgcgttg agttgggttaa ggggtggttca cgttgtgatt ggcggaagtg gttgggttcaa 60
ctccttctac cattgggtgag ggaagttgaa tattacattg taaaaaaatc ttacatagtg 120
tcgcctttcc ctgtgaaata ttnttcgcaa tagaaactaa tcttctgcta gaattatgag 180
tgaaaaactaa ccaattaaat atttatacaa ttaatactta cagtatttct cataattaaa 240
taaactcatga tgatcaaata tatattctct tctaaaagag aatcaattca acatacacia 300
gtctacagga aaactatatt atgagtaaac tatca 335

<210> 625

<211> 514

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 625

ntganncccc tgttgaaccc tatagtatnt accngtcact atttcgtact caagcttatg 60
cttctaccgg ttntggacta ccaaattgtc aagttgtttt tgcattgaaca aaacaatcat 120
attcatctca caaaaaaaaaa tcgattccaa atcgatgttg ttggctaaag ataattgacat 180
ctcttaactc cttggcaatt gtatcaattc gttcaagtaa tagtttaaac gataaaatcg 240
agtgtgagtc cacaagaact ttgactgtac tcagagttaa tatatatcca attttaagta 300
ataaattaat tgaattgaat atttggtgag tgatgacgca taaatataat tttgatctaa 360
attaaactac aaaacanagc atgtgcaagg gtgagaaaac aaacactcaa aacagtgaag 420
taacgattga tgcagaatta tgaacatgtt ggggctcagt cagcctacca gaactactct 480
ngatgcaaca ttaaggatnt ntctctatct aacg 514

<210> 626

<211> 314

<212> DNA

<213> Glycine max

<400> 626

gacgaagcgt gttgtgaatg gcgtctcgcg tggctagcat ggtatcgaca gcatcattgc 60
gtgatgacca cttgatgtaa ttggagatgg ccggaggagg cttaccatat gtgacctcgt 120
atggagtga gtcgggtgccg gagtgctgag acgtgttgta agaccattca gctagggcta 180

agaattaaac cagtatgccg ggttggtgtg aacgaaggaa cgaagatatt gctcaatcgt 240
accgctcatc acttcggttt gcccatcgga ctggtgatga tacgccgtac tcatgcgtat 300
cgctgtccca ctga 314

<210> 627
<211> 264
<212> DNA
<213> Glycine max

<400> 627

agctgtagga cgtgaaatca tgtgcagtca tgtatcttat agtcctctca cgggggggag 60
gttggtccat gctctcagaa tgtgcaaaat cagaatgctc agaatcagaa tcttcagaat 120
cagaacgctt aagattatta cgcttccaat cgagatggtc atgagcacca ataacatact 180
gcacagattc atcatgagcg gcctgctccg gatgaccaa aggaataaca tgatgcctaa 240
ctcatctatg aaatgtccta tcta 264

<210> 628
<211> 522
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 628

tggggaatga tacctnagt agtangcgtg ctattacctg tcactatcga atactcaagc 60
tgggagagga tgcttcaatg gaggaagaga tttatggaca gatttataga gggggtagca 120
cgaaattgaa ggaataatat acggagagaa gtggaacctt gaagtatgtc ttacatgact 180
ctcattcatc aaagtacaa catgtgttgc acatgcttct ctttatagac taggtagctg 240
gcttgagaag ccttggtgtg aaaactgtct tgataagctc atttgtgaag cattacgtgc 300
gatgctagag cttagctcca cacaccggg tgataactaa actcacctcc tggagaagct 360
cccttaacat gactcctaac gaagctatag cttaactaca catacctccc taatacttaa 420
gtcaccttc ttgtaatgag aagactttaa ctgagctach tggcccctgt tgtagctata 480
ctcacacctg tgacaaataa gcaatggtat gcaatacact cg 522

<210> 629

atcatagcan ataattaacc ttctaagagc tntccgcctt ttagcanagg ctgctttaga 240
 cctgatcgat gaggtacaac tcccttaaca aggttccgtc aatgtggtcg attgattatg 300
 gagttaattt tcaacaactt tgtagggata tgcatacatc atggatgatc atcatacaca 360
 tacta 365

<210> 632
 <211> 396
 <212> DNA
 <213> Glycine max

<400> 632

tagcttttgg ttttggtgaa cacaattaat ctattaccat gtaactgtaa togattactt 60
 ggggtgttgt gcgtgatgta atatgttaca tctctatgtc ttcacctcgt caaccactat 120
 tattactaca tttacttccg ccacccacct cagccaacta aaatgcctca gcctctcttt 180
 ataagccaac cttacctat gcacaaatca caccaacacc tgttctaaac ctctgtttac 240
 cacccttctc tgaatacaaa tatcttatgc tctcaaacc tctatctcct cttcaagatg 300
 acaacaatc ctttaagagc aaataatatg gaaaaatatt ccaagaagaa gcaaggagaa 360
 tgttcatagg atgttatcaa cccgttggac ttggaa 396

<210> 633
 <211> 517
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 633

gcgggaatga acccatgtga nacntgttg atacgatggc atctacgtga cactgtgtag 60
 tgctccagcc cgtgcgatta tgaggtaccc gtctcctgtg gtactaggtg gtattcgggc 120
 gagggtgcat aacatttctg acgatccacg gatgacctat aagcacccca tacgctagtg 180
 gcgccctatg gcagagctca ctatctcaca catgacacac atacgtgggtg gtgtaagcac 240
 cgggtacacc tcaagccttc tgagctttca catcatacaa gtaattcaac aattattgat 300
 ccgaactcac acatgatgag ctgaaggcgt aggcgcagaa ctctgctgga acacatacca 360
 gtctcccgac ttttcacatg caaatatccc atatgcattg gctatggctc acgacgttga 420

ccgtgtggat cgactagcat agactactgt gcgtatctag tacatattct acatcttgac 480
cgatgtgata tgctatagaa tggctagact cacgtag 517

<210> 634
<211> 390
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 634

cttgagctat aattcattgt ttgtggtatg agtctacatc aaacaattgt attattgatg 60
tttctgtcac aatcaagtga tagttgatgt ctccatatgt gcgtacactg tgattatggt 120
ttcgtttcta aaattcattn ggagtatcta ctgttgatgc tagatgagtg atccttcttg 180
atttacaatt attgtctcct aatcaatcga gtgttcacatc tattattggc tgctcatatt 240
ccaatcatgt ctagttaaaa tgcttgataa tctttcttgg tgttacttct aatacgaata 300
aagagagact tcgcctttga caatcacgtt aagatgtttt gagaggaaat acttgagtag 360
acatgatcat tgtacctata gacgaaagtc 390

<210> 635
<211> 252
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 635

gattgatagt tctaagaa taatcttata aagaatacct ttctcttag caatgaaaaa 60
gtgagtcctt attttgttnt ggacaacaca ccttctttg ttaaaggaaa catcaagtcc 120
attgtcacat aattgactta tgctaagcan nnatatgtta agctctttaa caaaaagtac 180
attatcaatg ggaggatagg gatcaatact tantctttca tactccaact atttctctt 240
ctatttctt cg 252

<210> 636
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 636

cagctccatt gatatcattt atttaatgca atgcacatgt ttcgaataag atttcttgta 60
 tcacaggaaa agatctaadc caagttgaat tacggatcat actgattgtg gtgcgaaaac 120
 tggcttgaag tagccatctg ggtaatatcc aaaccacgaa gtttcctctg gtattaaaac 180
 agtgtcgcgc tcacactgtt aatgtataaa ataagactac aaatatataa gctttcctca 240
 tgaaatcaat tagcaatatt ctattcataa tatcacaata atacatttag aggacttacc 300
 atgataagta ccanattctg caagctactc aatcttttct tgtaagaagc atttctctta 360
 tctggtattt cattgccatg cactggaaga aat 393

<210> 637
 <211> 391
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 637

gctgtctcta gatagacaat cttgccctcc tgaccttacg gtgacttacc actttgatnt 60
 ctttttttgt tgggtagtaa ctgtaccatt aaatataagc atcatataag aaactagctc 120
 acaagaaaaa tctggtaaag actacatatg annaaatagc gtgggtcttaa acaacaattg 180
 taacgtaata nactgaaccg ctgggtaatt tcttgaataa ataaattcag atcttcacaa 240
 tgataatttc ataaactctt actgtacaac acattgactg aagacaaaca ttatgcataa 300
 aactatagaa gcatggcata tgttcatctg caaccaacag gaatgtatgt cacacattga 360
 aacagatatg tnntatctca tactcattct t 391

<210> 638
 <211> 416
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 638

ctgcacaagg ctcttaatat ttgaagagta tccttgagga accttcaccc gacgaagaca 60
 ctaacaaaaa cttatcttct ccttcttggg caaagtatgg caggctgggg gcaagtaaatt 120
 tttcttccca tcagaccttg tatgcaactg tgatcgtata cccatatcag ctagatcttg 180
 acgggtattc aagccatcct tcgtcttggc ttgaatgtta aggagcgtcc caatgactct 240

atcacagaca tttttctcca catgcataac atcaatacaa tgtctaacgt caagatcaca 300
ccaatacggg agatcaaaga atatggacct cttcttccat atgcaactnt gactattatc 360
cttcttttga gtcttcccag atacagtatt cacgtgttca acccgataat atacct 416

<210> 639
<211> 298
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 639

gttcttgacc atattgaacc ccggtgacaa caattgcttc aaggacagag acaatgaaga 60
ggtcgctgga caacaggatc aacaaatgac ctatagtatg gttcttgact tagatcgaac 120
ctgaggacaa cgatattaac caacagacca cacaacaaat gaataacagg acaagagaca 180
atggagctgc acaacatagg gttaagagag acaaatgatg agggaataat ctcanatacc 240
ttgcaagcat ggtagaaaact ccaaaccacaa actgagcata ctcaaattat ccaatctt 298

<210> 640
<211> 405
<212> DNA
<213> Glycine max

<400> 640

atcactatac acattagatc tgagtaagat ctactattga agtactaaaa tgtacacaca 60
catacctcgc cacaaagagc aatatacttg gaaagcagct tggtgattct atcctgctca 120
tctttcttga gaattcgtag tttctcttca ctcacagatc gtgcaagaag attcccaact 180
gcaacaaaat cacaaaacta agctaaaacc aataaaattt caatgcacac acacaatcga 240
aattagagct gaaacataac tccaacttac atggtgtata gacataatcc ttggtactat 300
tgtgtgtgac atggagaaca gcaatggaaa ttggatgaga ctccacttct tgagagccca 360
attcatagtc ttaagtccat cttgcacatc attaccaaca gcaac 405

<210> 641
<211> 312
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 641

taattaagca atatggatga gatattctctc taattcagga tctaattata acatgatatg 60
atcaaagtga agaactaaca tcaactcaatt aagatatggg aattaaaatt acacagtatg 120
tgaatataaa gaaactcana ttgtatctaa actttctctt ctactaatat tactattaaa 180
taatatgttt taactatgcc tttaatgaaa gtgtgtttta actatatatt accaaagaaa 240
catagtgcaa cctatataaa acgtgtatnc gaacaaacct actatcagta agtatacttt 300
atacgttaat ct 312

<210> 642
<211> 482
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 642

tatgaccctt ganncccttga tccctgatta ctgcggntta agataaccgc angccgaagc 60
tgtatatgat ttatgaactg gccatcattg tccctatgca agtctngcac cttatgttgc 120
tggtcgacac aaccatcata ttccgactga gagtgtgtgc ttaaagaaca tcacgtcgaa 180
tcggccgcat cgctgtcacc gctcatatag agtaacatta tgtggactac ttatagcggc 240
gattacaggt ggggtggtctt ttcatatatc acatgctaca tccctttctc ctaaatgcaa 300
aagacttcgc tcgcttcata gcttatactt actctccttt gaangacaga cacatcacat 360
tgaacatcac caaaaatata ttttggcagg ctgccaaagac aaagacttat gctgtatttg 420
atgaccataa cttaataggt cacttccccg agacctagag aaaccgatct tcgacttaaa 480
cc 482

<210> 643
<211> 514
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 643

tggggatgac nccattagan ancncgtgat ttgcngacac tatacagtac tcaagctntg 60
agaaattatg gctgaggcta agacatgttg tgggcccatt gttcaaattg aggcatttat 120

tncaaagtat

310

<210> 646
<211> 297
<212> DNA
<213> Glycine max

<400> 646

agcttatcga gatccgtgat ggataggcaa tgtttcggca taatgtggaa cctgacgttg 60
ttgatgtaca catgcaaadc ggagaagatg gagagttatt tgatggcaaa cacaatcatg 120
ttgttcagct tcacgacttc cgcgtgcac accactaagg tgaggatgtt gaagccattg 180
ttgcggataa gaagcatcac gtcaatcaac atcaagaata ccttcgcaa ggtgttggtc 240
ttgttccaat gaagcgtatc agagggttaa gtgatgcaat ggcgaagtgg gttcgag 297

<210> 647
<211> 372
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 647

ctaactgngc agtantccaa aaatgtgagt gacatcacct tgagggtgggc aacaaattaa 60
gtgccttcac cataatagca cctgcacctg aacctatgaa gcttcattct acacctgaga 120
aagatgatga gatatttcct cataagagaa agcttccaac gcacatatta cgcaactgac 180
ccttttttcc tctcatgact taaccacccc tctacctct attattccca ataccatata 240
tcaactaccc gtacatcacc attctgcacc taccttacct ctcccttgag acaaactctt 300
tatcgcttc acatgtaacc ccgtaaaccac tacaatctct aatatatgat tgctcaactc 360
tcttcaacct ct 372

<210> 648
<211> 375
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 648

tttgagccgt cgacgacng ngatccttag agtcacctga ggctgctgct tacatcaaca 60

cttcaggggc tgtactactt acatggattg atgggcctat gcagttgaaa gcctggagaa 120
agaggatgcc tatngttggt gggatgaatt ctccagattt acctgggtaa actctatcag 180
agagaatcaa aaccttgagt attcaagagc tgagtctaag acttcaaaga gagaaagact 240
gtgtcatcaa gagaatcagg agtgaccatg gcagagaatt tgaaacagca ggttcactga 300
atgtgcacat ctgaggcata ctcatgagtt tttgcagcat tacacaaaca gaatgggata 360
gtgagaggaa aacag 375

<210> 649
<211> 234
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 649

tttgagcgctc tcgatatatg acgagactac atcatacatc ttagtaanaa gttatagtcg 60
tttgaatatg ctgagagctt caacattcaa ttacgagcat ctgctatat tacgggactc 120
aatcagacat ccgagtaaac agtttgttgt ttgaattgct tgagagcact cacattcaca 180
ttctagcgcc tcgatatata tatggactct atcacacatc cgagtataaa gtta 234

<210> 650
<211> 407
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 650

tgcaagcttc tttgagaaaa cttccttgag aagctagtgc ttagctacac acaccctct 60
cataactaag ctacctcct tgagaagctt ctctaagaag attcctaaag aagctagagc 120
ttagctacac atacctctct aatagctaag ctacctcct tgagatgaga agctagagct 180
tagctacaca ccnctataa tagctaagct ccccccatg acaaaaaaaaa catgacaata 240
caaaaaaaaa agtccttact acaaagacta ctcaaatgc cccgaaatac aaggctaaaa 300
ccctatacta ttagaatggc caaaatacaa ggcccaaacy aagaaaaaac ctattctaata 360
atttacaag ataagcgggt catgcttagc ccatgggctc gaaatct 407

<210> 651

<211> 337
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 651

cgcctaacta aatatgatat tgagtgtttt ttacatgcat ntacttaaca agaaacttan 60
 acattagnta agttttaaact atatggttgg aaaaagaaca agtggaagag tgaagggttc 120
 caaacaacaa tccttgacaa aaaataaaaa aaagctgggt ctagcaaata aatcatatca 180
 cactaagaat gagaaataat accagcatcc ctatccaaaa aatattcaag gatgagacgt 240
 gtaaaaggtc acgaatttca tgctgctaata gatatatcta acantaataa tctngtcctt 300
 ttataccttt tanggtatta tggagatgga cgggagt 337

<210> 652
 <211> 178
 <212> DNA
 <213> Glycine max

<400> 652

cccatcatga tgctctttct gaacagaaaag tacctgtgaa cgtgcactaa attgcctatc 60
 aactcactac gagagaaatt gaacttatct taaactctgt gatatgagta tcattacaat 120
 tatgaggcat ctacgtgcat aaaatggatc aggttggtcc ttatgacctt gctgccta 178

<210> 653
 <211> 354
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 653

tagctacaca caccatcta anaactaagc tcacctcctt gagaaacttc cttgagaagc 60
 tagaagctag ctacacanc cttataatagc taagctcacc cncatgacaa anaaacatga 120
 naatacanaa aanatcctac taaaaagact actcanaatg ccctgaaata caaggctaen 180
 accctatact actagaatgg ccaaaatata aggcccagac gaagganata cctattctaa 240
 tatntacaaa gataagcggg ctcatactta gcccatgggc tcgaaatcta ccctaaggct 300
 catgagaacn ctanggtctt cctttggatc tctagcccaa tctacttgga gtct 354

<210> 654
 <211> 445
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 654

taatagggta agcactacaa catttataat ttcaaagcaa agcacataat aatagctatt 60
 gtacataggt ttgatcacac ataacacact atagaatatt attcgtcata attaacaaac 120
 aattctaaaa gctatagcgt catggtactt taagatgaca tatagagttt taggtttatc 180
 aaagaatttc ccataacaaa tccaggacta tatcccaaca ttgcaaatac tacaagagca 240
 agcaatcaat atactttaga atctcacacc catattccat ttcaagcatt atattttttc 300
 atatttagac ataacttggc ttatgttcta ggccaatatt ggactttatt ttacaccagc 360
 ttattggact ntaagaatac atcccgcaac atcaaatgac tcaaattccat tactgcaaaa 420
 taaaatacc cagaggaact acaac 445

<210> 655
 <211> 520
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 655

atgtaattga aacctttgta agncccgta ctatagaaca ctcaagcttc taagattatt 60
 aataatatat ccattattca atattttcaa ttaattataa aaacaaaatt atttatctaa 120
 aaaagaanaa ttaaaaaaat attttgaaaa caaaataatt taaaattact aagagaaaga 180
 gcaactaaga ttttgacaga aaaaatgaat gcaaaaataa cacaattaaa attaaaaaat 240
 aataaccatt aatgtcttac atttttatgc ataaacatat atattacttt taatttaaaa 300
 ataaaaatat tttagtcatt tgtgtgaaat taaattactt acaacaaata aatttaattc 360
 aattctttta tagtaaaact ctttatatat atatatatat atatatatat atatatatat 420
 atatatatat atatatatat atatatatat atatatacac acgtatcagg gacatatgtg 480
 ggataatatg actacgttat atatgcgagt ggagatgctg 520

<210> 656

<211> 420
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 656

agtacaataa gctntatggt ctaaactctac tggaagatga catcccttgc catgcaccat 60
 ctagaatggg gaaagactgg taggggtctt ataggctatt ctatacacc acaaagcatc 120
 atccaattta gcaaaccaat ctttactaga attctcttta tctttgtaaa tgagacttca 180
 acttgaccat ttttttgagg gtggtaaggc gatgcaacct tctgtctgac attatagtgc 240
 tccaatgcct tctgtagttg cacattgcaa aaatgggaac cccattact gatgagaact 300
 ctaggagttt ccaaccacg aaaaatatc ctcttcacaa accgaatgac tatttttgca 360
 tcacctttct gtgtggcaat ggcttacacc cattttgata gaaactcttt aaatgacaca 420

<210> 657
 <211> 265
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 657

gtatccattc agagaaatnt ccttttggga ggtcatcatg aggccaacaa gattccttgt 60
 gtgaagtng acacagtttg cctttctaan aataaagggg gccttgngat taaagatttg 120
 tctaaattta atgaggctnt acttgacana atgggggtgg agctggctaa taattagaac 180
 caactntggg caagaatctt aatctccaaa tatgggtggt ggaaggagtt gatctctggt 240
 ggaaagagca nattntcctc tcata 265

<210> 658
 <211> 360
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 658

tgaccttgtt tacagcgacg ggcaactctt agaactgaca gaggcccata atcagagctg 60
 gaaacgccag gaccctattg gacttcttcg agtccactgg gtgtcttatg ggtgcgatcc 120
 ctacaaacaa tagatgacat caganatcag ttgagcgatg tgcatactta cctatgtcac 180

gatggcatga ccttgctggg ggcacgggca ccctgtaaga ctgacagagg cccgtaacct 240
gagctggaaa ccctagggcc ttgttgggtt tctctgagac cacggcgtgt cttgtgggcg 300
cgatccctag caatagtgga tggcatcaca aatcaactga accatatgca tacttaccta 360

<210> 659
<211> 360
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 659

ttatctagta cattgtnttt gaaacaagca ccctacagta gaattgccaa caatagacat 60
tgtttcagtt agggttttca attatcatta ttgacgactt ttagtgagcc tcctaaagaa 120
tgtaaacgcg tggcttatgc caccctcttg ataaactaag aagaagaagg tgaataataa 180
ataatctttt ctattaaaat aatacggtag ttggtagaag gtatttataa cattaaatag 240
tactctcttc atttgtaaac gatatttatn tatatacaca tattaagaac gtcaaaggat 300
atatattaca taataagggt tcattgagtg ctatagattt taatattact ttagttttat 360

<210> 660
<211> 390
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 660

agcttgtagg gttatagtct cacgattgtc acgtgctcat gcaacaattg ttagccgtgg 60
ctatacgaga catcttgcca aacaaagtca gggtcacgat aacttgccctg tgctttttct 120
tacatgctat gtgtagcaaa gtgattgatc cagtaatgtt tgatgagttg gaaaacgaga 180
ccgcaattat actatgccag ttggagatgt attttcccc tgctttcttt gacatcatga 240
ttcacttgat tgtgcatctg gtcagagaaa tcaaatgtcg cggtcctgtt tatctacgac 300
ggatgtaccc ggctgagcga tacatgaaca tcttatnaga gtatacaaag aatctatatc 360
atccgaaagc atctattggt gagaggtaca 390

<210> 661
<211> 111

<212> DNA
<213> Glycine max

<400> 661

agaatgaagt ccactcaaac ctgaaatctc caacttccac tcgtagacac gcacttcacg 60
actaccgaaa tgccctcctt ttgcgatctg gagcggaaat gatggccaaa g 111

<210> 662
<211> 404
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 662

agcttgtacc agccactana ccttcaattt caattgtang ctntcctaata ttcttgacct 60
tcttctacg agagaccaac ttcaactcag tatgttctg tanataatta taattaatta 120
gaataaataa aattgtatat aaaatntaaa tatgttataa ttaaagaaat aactacctct 180
cttgcccaca ctttggttac cacatgatta acatatgatg tcaacactaa tgtatcttgn 240
ggcccacctg gaaaacccta tgaatcaaca cctacatcct ttgtaattgg atcatgaggt 300
tcttcatgag tctcatcagc agcatcatcg atatgcccac tatcctcgac aatagttgca 360
agtgttcatt atctacgtgt cgacactttc aatcttcgac gctg 404

<210> 663
<211> 319
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 663

gcacgagaca tcagggttta gtattccaag atacaacata tatcgcatga acattgttag 60
atagagaaca tccttaataa catcagtcac ccagtaggaa gaccaacacg ttctttatct 120
gtcttcatac accactactc acgtgattta attttggata gtttagttgc atacttgtcc 180
ataccacgca ccanactntc atccaaaggc acttattttac tgaaccacag cntaccaag 240
tacaacagaa tgctcgggag ttggatactc agtattcact taccgggtta tactactttc 300
gtgatccagt gcacttgtc 319

<210> 664
 <211> 349
 <212> DNA
 <213> Glycine max

<400> 664

aagcttctat agaatgttcg ttcctaattt ctctacaatt gcatcacctc tcaatgagct 60
 ggtgaagaac gatgtggcat ttacctgagg tgaaaaacaa gagctagcct ttggtttgat 120
 caaagaaaag cttactaagg cacatgttct aactcttctt gacttttcta aaacttttga 180
 gctacaatga gatgcctttg gagcgggagt tggagctgta ttgatacaag gcgggcacct 240
 tatttcttat attagtgaag gacttcatag tgccaccctc aactacccca cctatgataa 300
 agagctatat gccttaataa gagccctcca aacttgggaa cattacctt 349

<210> 665
 <211> 295
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 665

gtcacattgt caaagctcca atcttgnnt tgaaatagag agagcaacaa cagcaaanac 60
 tacaatcgaa gaaaataaat cataatcgaa gagcanaaaa aatatacagaa acaaggttct 120
 aacgtttctc tcacacaang ccttttattt cctctnctaa tccatttttc ttttcttttg 180
 gtattccacc anagacaatt tttttctaag ggaaaaaaca ctcgactggc agagaaatag 240
 tgaagtgaag agagagactg agagaanaga tattattctg gtgacgctga tgtgt 295

<210> 666
 <211> 429
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 666

agcttggcag tgtgacattt tcttctgagt atttaatgca gcagattaat gaggtttcat 60
 tcttatcact gaatttccca tcgaagcaca aacacaatca ggagcactag gtttaataata 120
 atccttagct tgagatgcca atttcctctg ctccctgcgt agttatatgt caaaagtggc 180
 aaagactctg catcatcata acaatcaaac caaataatgt gcataaggat ttagctcttc 240

tagagtgagc aagtgtgtaa agtaagaaag taaaagtgaa tgtactgttg atttgaattg 300
 atgtacctgg tggccctaac taactttaaa acaatgggga aacaccctta gtttctcact 360
 ttatacattt actaacttta gaggagtcaa atccctgcat cagaatatga acaagaaaac 420
 atcacctta 429

<210> 667
 <211> 459
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 667

ntatcaccag gatgatcaaa atgtttttca tcattgcttg agttggaagg ttgggaatgg 60
 ggataaagtc agcttttggga aagataagtg gttaggggaa ggctctactc tacaacagaa 120
 atacaatcag ctgtttctaa ttaatagaca gcaatctgac cttatttcaa tgatgggcta 180
 tttctctcat gatacatgga gatgggactt gaaatggaga aggaacctgt ttgaccatga 240
 aagtgatcta gctgtcgatt ttatggaaga aataagctct tttcattttc agagaaatgt 300
 taaagacatc atgacctgga aagctgatcc tagtggtgtc tattccacga ggtcagcata 360
 caaattcatg ataacccctt ctttcccagc ctttgatctg agatcctcaa ctntattatg 420
 gaaattgaag attccccaga agctgcagtt ttcacttgg 459

<210> 668
 <211> 306
 <212> DNA
 <213> Glycine max
 <400> 668

ttgctgtgat gaacaaaatt tagccaatat tatcagataa atcaaatact taccgtagaa 60
 ttactgaaaa tagaatctac atcatcaaga ctaatattag cctgttcaaa tacaggaggc 120
 ttatatccct tcttgaacca gtcattctca atgacctcag taaatgtaat ccgctgtgtc 180
 atattattaa aagaaaaaca gatgccggtt gttaataata atatatttgg tgaaatggag 240
 ggaaaagtaa aatattctgt atacactaga ttagactaga gtactatata attaaacaaa 300
 gaaaac 306

<210> 669
 <211> 363
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 669

gagagtcac ttgatatgac agctntggaa gtcctcttac gagactatgc tctagcatga 60
 ccaagcttct tatgccatac ccagtgatgc tctntgattt anagtaagca tgagaacctt 120
 tgactggaca gatcaccaag tttaatatata taaagaattc cttgtctctt agcaaagaag 180
 agtgaagagt tgtccttgnt ctgaacaata cacatccttg ttaaagggtga cattgtatcc 240
 actatcacat aatcgactta ttctcaacag aatatgcttc aatcctttaa cangtaanac 300
 attatctata taagggtagg gaggaacaca tactttacct acacanngta tcataccttt 360
 ctg 363

<210> 670
 <211> 505
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 670

atatgaatcc tgcangcagt natcccttag aatgcngccg gtcgnancct nggaccgatt 60
 catgcagaca agggacncct gtanagcata cataggtttt tgtcagctga actaatgagg 120
 cgaacatgct catcatgctg acgcaactca actcatgggg ctctctgaga taatgataat 180
 taagcagaaa cctaagatgg cctgtcaatc ctatacagca ataattacag aaccctcct 240
 tgccttgccc accctttggc taaccatgag taacacacga cgtcaacact aacgtatgct 300
 gtgcccacac ctgaacaact ctactaatca ccacactacc tcctttggaa ttcgatcatg 360
 aggtttctca tgagcctcat caggggaatg attgatatcg cccattattc ttgacaatag 420
 gtgccagtgn gcattatcta cgtgtcccca ctctcaatca ttnacgctg agggcgttct 480
 gctcgacca ctaacctntc taccn 505

<210> 671
 <211> 228
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 671

cttattatta ttccaataac tntgtgtgcg gaacaataaa aataatactc ataaacatat 60
tananagcat ttaacaatga ggaanaaaat gtcataacc atanaagaaa aaaagccaca 120
agaaaataaa aataaaatag tatntattnt ctanactac ctnccttatta cctaattagc 180
tcaatcttgc aaatttaaaa tgcacaattg accaaaaaat aacttgat 228

<210> 672

<211> 226

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 672

agcttgagat gaggaagtgt tgaagggtga attttcctgc ttttattggt gaccacagag 60
tggtacctgn agatatgtcg cggnggtcag gagaccttgg ggacgtcagg tgggggtgcta 120
ttgccccaaa ccaagcttga ccaatcccg cccaaccccg gcatagtcgg tcagtggagaa 180
catgtgacgt acctaagcag gcgagcttct tgcagtcaca gataaa 226

<210> 673

<211> 296

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 673

caggcaacta actcctcttt canaaccatg ctatgtgctc gcgactgggtc cctttcttcc 60
tttcgcaact tgagttcact attgctaccn catagagctc cgcgaaattt gttccggcca 120
tactcttctt tgcgagccct cttggtctct tgttcaaggg ctcttgccgt aattgcattc 180
tcttcccgta acccggcaca ctcttccga acgtgtgtag cggccaactt gaacttcttc 240
ttggcaagtt ttgcctttcc taactcgcta ttgagagctt ggacttcttc gtgctc 296

<210> 674

<211> 113

<212> DNA

<213> Glycine max

<223> unsure at all n locations
<400> 674

gctgcaagct tggattgatt cggctctgaca agggattaat gnttagtaat ttaggctaca 60
acattgaaca aaagaggcaa atatgatcat catgctttga taaaaaaaaa atg 113

<210> 675
<211> 320
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 675

atgagaatgg ggtagatttg gagcatactc tcatctcaaa caagtctata acattaatct 60
aaactcgctc aaactggttt tacaacgaan aatctaccga atcaataatt gactcctcaa 120
caccgaattt accctagaaa tggctcttgc cttcactttg gtcactcatn ttctcattt 180
gctcagccca agctgtccca taagtcctaa atgacgattc anactaggat taactcactg 240
taatattcaa ttaccactaa atccagaatt agctnttcag acccctcaag cattacactg 300
tgtcactcat atcactacat 320

<210> 676
<211> 348
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 676

aaaaaaataa tttaatcgaa ctgatcgaaa ttcggaacat cactctaate tgtgtcaaat 60
acaanaaata aaaaaagaac accatatatt taaaaaaaaa aaaaaatttc tccacgggac 120
ctatgcaaca ggatagcaaa aaattcagtg cgctgagtga ataaatgctt ccaaaagggc 180
gtaaatcgag gggaacgaaa caaaaatgat tcatatctga aaaaaaaca gacaccaa 240
accaatgaaa gctttgcaaa aaaagcatag atcagcgcat atcaaacata gagcatgatt 300
acataatcag atgaaaacgc ggattatata tacaatagta atatgagt 348

<210> 677
<211> 312
<212> DNA

<213> Glycine max

<400> 677

ctcctcatac agtgaaactg gcacaagtga ccagacatct gatacggtag aggggtgattt 60
attattcatg tttcatagtt gtatattata atcgcacgta tatctataag aatgatgatg 120
ggtagcactt catcaagtga aacctctatt cttaacacat gaggaagttc acagtacaag 180
tgaaccattt ggtattctca atgacaataa gttaaaaata tcagcattca atgtagagag 240
ctcatccgaa gaagtccggg agatcttcat ggggtgcatga gctacagatt ctgcaagagt 300
tattgaagtc ac 312

<210> 678

<211> 381

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 678

cgctctgnca ttcagnanag aaggagggtt gagtcccggg ttatggtttg atcccacatc 60
aggaacgtta gcaatatcag tacttgggtct tccgtatgag tttgaggagt caactgaggg 120
tatagtctca atggccagct tgggttatagt atcttcttgc gtacctgatt ctggagagcc 180
ctgcttctgc ataaaattgt cgacagtatt atttctaagt tctgggtggca tggattcacg 240
ctcaccagat acagggattt ggttgatttc aataactgag tgctttgtgt catcagcact 300
caacttattc agctgataga cagaaggcat agactgctgt ccagatggga gcctcttctc 360
agtgtttcaa ctgatggact g 381

<210> 679

<211> 381

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 679

ctaaacttaa atctcgcat actaataaat ggttaatact gtattattct ggttttacat 60
ttntgttcta tgtggcttct cagattatct cttcacctag taaataagag actaatctac 120
caagcatgat ctggtggtga naaagatcaa ccctctctag tctcaatggt acctttctca 180

caaacaaagt tcagaggatt gaatcanaaa tcacattcac atgctcaagg gatgtgagcc 240
 gctgtcactc acatcaatgt atggtatgtc tgtaatacgt tggttgagtt atgttattgg 300
 caaaatggat tggtcacaac ataaccacta tcatatggca cataactcac ctattgagag 360
 catgatcaac accttcagta t 381

<210> 680
 <211> 317
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 680

agctngaccg atccccgacc aaccgcgaa nagtttcac gtgagtatga aaagaggcac 60
 atgtaatcat cctgctcgaa caaatgacaa cactggggca aataaagagg gtgagaatgt 120
 agaacaaacc catgctgcga ctgccattac tatacggaca aggttcccac caacccaaca 180
 atgtcattgc tcaaccaata acaacccttc tccctaccta ccaccaggt aatcacaaag 240
 gccatcccta aatcaaccac aaaaccatc ttccacacaa ccaatgctaa gcaccacctt 300
 tagcacaac caaaaca 317

<210> 681
 <211> 151
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 681

cttgcaactg aacgccaaatt gttatcatca tataactaatg gatcacatcc agacagctta 60
 tccgaagggtg ttctaattgt gtcggaaatn ttgcacagac tacaaaaactc tggcgattaa 120
 tatttctgac gaccattatt ctaacggatt t 151

<210> 682
 <211> 334
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 682

acatgataca tgtcttggtt tggctgggct caaagataaa ggagatgccc cacattattt 60

ccatgacaca natgcanaga tgatgatttg gaaacttcat gcaaaaactgg tcatgcatgc 120
 acctatgtgg aactcaagt gtcaaaacttt tatggatcatg tgatgctagg gctcangatt 180
 tagatcaacc caatgttgcc aaaatatgtt cttttatcca tttgtgcatt catccgagtc 240
 catttcgggc gttcgggtgaa atttcacagt gttcaccctt caggtgtaga cacattnttt 300
 ttcaaaaact agttatgatc aatgaatttt tttc 334

<210> 683
 <211> 407
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 683

gacaactctg atatcaagag acttacacac acacactntn tcctagtcga tcaactcacat 60
 anatntccat tctcnccctt tggttttgag tttatgcntt catttgaaat tagttaatta 120
 cttatgtgag ttcttgattt attccctata tctctcccc tttggcatca acaaaaagcc 180
 aaagtgtgta acaagtataa gacacacata tactattaat cattcacaag gcatacattg 240
 aagaatataa accaatcatg aagcagcaaa catgaataga tcanatatat aataaccaca 300
 tagtcatata atataattca taattgttca ttcacaccat gccaatatag aaaatactaa 360
 atatccaaat gtcataataa tatatggtat ttggataagt cactaca 407

<210> 684
 <211> 278
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 684

agcttccttc anaaattacc atatttgcgt ggtgactact aagagatagg ctaccaacaa 60
 gatcaaattt gaggagaaag caaattgagg tcgaagatcc actgtgtcca ttctgcattc 120
 agctgaggag agcgcttgcc aactattctt tcagttggag aaggatactc cgctatggtg 180
 ggaatcatta tcatgggtga angtagtggg ggcttgtcca aatcatacaa ggcaacactt 240
 ncttcaacac atatatggag cgacagatgg aatgagag 278

<210> 685
 <211> 454
 <212> DNA
 <213> Glycine max

<400> 685

cagagggtgt ggttgtgtta tgttatcgat ttctactctc tattcatatg tttccgtatg 60
 taattatgaa aacttcaactt aatttatgct tagttttgat tcttattttt ggctgtgttg 120
 tttatttggg gactgggtgc cattaatatg cttttggcag tgctttttca tttctaacac 180
 gttcttgtgt tctgcaatct ttttgcattg aatttcttat atacttagta gttggaatgg 240
 gcactgtgtt atcttttcag gtctaacacg ttcttttgtt ctgcaatatt attgcacatg 300
 atttcttaaa taccgagtag ttggtagata ttgtttcttt taacagcata catactgcca 360
 tgtgccatag agatatagaa ctgcttgact cctgggtgtgc acaatgcatg tataactcaa 420
 ctactgtgc ttgaatctgc ggacttctga aacg 454

<210> 686
 <211> 304
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 686

agctntgctg atttagttnt cacttacgaa aggttcaaag tgagtctgaa aagaggcaaa 60
 tttaatcatc ctgcttgaac aaatgagaaa actggggcaa ataaagaggg tgagaatgta 120
 gaagaaaccc atgctgcatc tgccattcct atacggncaa gtttcccacc aaccaacaa 180
 tgtcattgct caaccaataa caacccttct tcttaacctac caccagtta tccacanagg 240
 gccatcctaa atcaaccaca aaaccatct tccacacaac ccatgctaag caccaccttt 300
 agca 304

<210> 687
 <211> 314
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 687

gcttctctgg gacctangca naccttcaac tcctccttca cgatcaaagt gtctactcgt 60

cattgggtcgc tntcctccct ccggagctta agctcgctat ctccagttcgg cattntcctt 120
 ttggatctta agagttgctg attngaacct ttattttgac cggtgggctt gctcgagtcc 180
 tgccctaagg gactacacct cttcatcttc ctccggtgcc tcaacttcct ccccttttgc 240
 gtgtgggatt tcagccactt acggtagcct ccaatgggcc cggtgtctnt gtctttcttt 300
 gcattatttc ccat 314

<210> 688
 <211> 120
 <212> DNA
 <213> Glycine max

<400> 688

agcttgacca atcccgaccc aaccgggca tattcgggtca gtgagaacat gtgacgtacc 60
 taagcaggcg agtcctggc agtcaacaga taaaaggaaa acacgaccac agagcaggga 120

<210> 689
 <211> 224
 <212> DNA
 <213> Glycine max

<400> 689

cttcaccttc ttgtcttcaa cgtgaactat gaccattgtt ctatcttccc gcgatgcttc 60
 ttttcatgtc cgctgagtg ggcttatagc ctaaaccata ctctccacga ttgcttgtg 120
 tatttatcag gctagttagt ccgacgttgt atttgctata cccatcctgg gttcataacc 180
 gttccccaac ataactcggg ccatcattac cgctgcatcg gaca 224

<210> 690
 <211> 209
 <212> DNA
 <213> Glycine max

<400> 690

tctgtgtggt ggtcggcaga ggagcataaa ccacacagtc tggcgacagg tgcagatatt 60
 tgagtcattg ccagttgggt taccagggtt accaaggcat ctagtattacc ttcaagcttc 120
 ttagtctcac ctgatgaaga tgaattcatg gctacttcaa gactcctct aatgacaata 180
 acatcatttc tggcactgaa ttgctggga 209

<210> 691
 <211> 386
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 691

accaagtggc ctagaatttg atcctggctg tcgacactct ttaagtgaag agccgcacgg 60
 ttggggagcc tatcctgtgc cagcatgcaa gcttaaaggg ctgttatgta agggagcggg 120
 ttacgcatgt aatataaaac cattgggtgtg ggattagcct acaactaatg ctattctagt 180
 tctctagata ggtgggtgac aggcattgga gacggcgaaa aagttaacta taataccgca 240
 tcatactaata gaaaacttcg catgctcaat gaatgcttaa taagatatgt ggctgcacaa 300
 gaaaagtgac acttacgang accgcgtcta tagatcgact cttgccggat ccttgcatgt 360
 cgtctttata acgcatacat tctgac 386

<210> 692
 <211> 187
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 692

gcttatattg taattgttta agttgtaaga ctctcgaaac accttgtna tcttgagnaa 60
 aaaaagatta ngtgcttaat ttgtatatct gtctataaga cattaaggct agtttatgtg 120
 catacaaaca tcaacaactc tacntaattg ttagagccag aaatggctta atagtcaaag 180
 aatactt 187

<210> 693
 <211> 614
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 693

tcgacacttc anttctccat tgtatactcn cganatcata attcntntac tggctactga 60
 cacncaaach annacccgaa gaganaattg atgcgtctcg tacnccgat cctctaagtc 120

tacctgccgc atgccagctt gtggaagagg caaataacat attatgcgtg ttgctatata 180
 actatctccc atctctcaat caatcctttg tatagtgtag gcttgtagac cacaaacttg 240
 catatgggga gtttgaggag gtgtacaact ctatctttac gctgtgagaa ttgttcatat 300
 tggctgtggc tatectttgt gaacatgacg gtttacgtgt tgggcgatgg tgtgcgtgac 360
 tcaccacatt tgctaaactt gtgaatgttc tctgatgtga cgatcaagat tacgattggt 420
 gctgcataac gggttctgaa ggatgtatac tgtacgggtg ntgactctta tctactctat 480
 tccacagcac gcagcaccca agctacttgt cctccggaac cataaatcgt acgacgtatc 540
 atacgttccg attgatcaat cgaatgtaac tatccacatc tcggttcgat catacctatc 600
 accctaacta tacg 614

<210> 694
 <211> 436
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 694

caagatgatg ggctcaacat atnaacaaac ccgctggcca atcatggagg ngggcggata 60
 atgccataga tgtgagtagg ncgcacccggc ccatactttt gaaggatgga acgacctcca 120
 gaaggtttat ctacgaagcc ctgtcaaaga cggtatgatt tcccgtgcca gagaacatct 180
 ttgctaattgc attccgttgt actgcatgac atggaaacat gctcggcgga tagagatcta 240
 ttgcacaaaa tgatacaaca cggctagcta gatgtcgtct atgatgggga ggaagaatca 300
 catgtttacat cgccgctacc aatgatcaag tgctcataat cctaatectt ggtaatcatt 360
 acctagacac tattccaatg acccaacacc tttgcatgat ggcagcaaca ttctattctt 420
 acttatcgca caactc 436

<210> 695
 <211> 425
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 695

agatgcatgc ccatctatta aatgttgcca gtataatctg ctctattatt ttacaacacc 60

cttgttttaaa gtgtttcggg aattatatat aatgaaatgg aaactttatg cttctatatatt 120
 tatgattcta tcgagttcaa ctgactatat accaatatgtt ttgtggtata aatgcactct 180
 aaccatctta attattacta aaaaaatgac atctaagaca gattgattag aatatgtttc 240
 atcaaattat ttgtaaaaaa atatatagtt taaaacgatt cttagaaaaa ttatttcaaa 300
 aaaattatgtt ttgtaaaatg gtttttaaga aaatcgtatt agaactctta aatctgttaa 360
 ctttttttgg tttaaaaaag acattataaa acnggttctc taaaaatcgt cttagaaaat 420
 ctatt 425

<210> 696
 <211> 532
 <212> DNA
 <213> Glycine max

<400> 696

aacattcatc tccatgtact attagtgtag caacatgata ataatccacc cagcacgtac 60
 atgatgcttc gtctcggatc ttatatcacc tgacgcatca agcttgagcg aaaagtgtga 120
 agatcacact tctactttt attcgcgacc acgagtggac ctggagatat gtcgccgggg 180
 tcaagagatc ttggggaccg caggtggggg gctatcttcc aaaccaagct tgaccagtcc 240
 cgaccaaac cgggcatagt ctgtcagtga gaacctgtga cgtacctaaa caggcgagct 300
 cctgcactca accaataaag aataatgacc caatgcagga cgctgcgtgg tgctggctac 360
 tatggtcttg gtgatactgg atatggctta gctaagatac ctcggtatga atcgatacaa 420
 gctaaaatga aaccgcagct atattccttg gtatccaacc acggtgggtat gttcaccttc 480
 tataatggat ccgagttgca tctcttgtcc tcttctactg gacggcgta tc 532

<210> 697
 <211> 455
 <212> DNA
 <213> Glycine max

<400> 697

ttataccatg gatgaataac agcgggtcag ctacatcaaa aactaccaa agaaacttag 60
 agttgacaag tattgcagct tacaagttc attggatact ggaacaaaca aaggctcgac 120
 taaaggaaaa agagtcattt taccttcaac ctttggtggg agcccatgtt acatggatca 180

actttatattt gatggtatgg caatatgtgg tcatgttggg ttcttaaate tttttataac 240
tctaacatgt tatccaaatt gtcctaaaat tcgtagatta ctttcacctt tgaatcttaa 300
accaacagac aggccagaca ttgtctcatg aattttcaga ttgaaatatg aacaaatgct 360
ttctgactta ccaaagcatc agctgctcag aaaagttggt gttgcgcgta agtttagaat 420
gatctttgct gttgaacgta gaaatcaatt gatca 455

<210> 698
<211> 431
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 698

gcttgtaaga attgcaagat catcttcctt gactacagtt ngaaaaagat cgccaccaat 60
ataaagagat gacaatttag agagtgatcc aagactttca aatggatctc cactgaattt 120
attagtagac agatcaagat atcttaatga tgaatgcttt ccacacgata taagaagagc 180
accacctttt gtgtgttgg aagaatctat ccgctcaata tttttaaatg cacctatata 240
atctgtcaga tggcctgaaa gtcgtcaact ctgaacttga agtcttgtga gtccatggga 300
aatacattga gcaagaatct ctaacagttc attaacctgc tcggttgagt ttgagattcg 360
agatatcaat gtcccttaag tcgcagagat tacccaaaga agttggaatg ttttcttcat 420
gtcgatacct g 431

<210> 699
<211> 381
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 699

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ccttttttga agtgacatca tggcagtttag gtccagctt tccatcgtgg attcagtcac 120
aaaacaaact tcaatatggt ggactgtcta acacgngat ttagattct attcccactt 180
ggttctggga accacactct cagggtttgt atttaaacct ctctcataat catatccatg 240
gtgagcttgt gactacaata aaaaatccaa tatctatcca aactgttgat ctaagcacac 300

atcacttatg tggtaaatac cctatctatc anatgatgtg tatgggttaa acctttcgac 360
caattcattc tctgaatcca t 381

<210> 700
<211> 389
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 700

agctntgagc canaatcatg actcaccata taccttgacc cagggtgaga atgccaatcc 60
ttaccctcgg aagcaaaaaa aagaagagaa cgaaaatttc caatcaaagg aaaaaggaga 120
aggaaaattt ccaatcaaag aggaagcaaa aaaggagaga aggaacattt ccaatcaaag 180
gaaaaagaga ggaaaggaaa ttctcaatca aagagtgcga gacagcaaaa agaaaagaaa 240
gataattccc aatcaaagaa tgggagatag aataaaagag agaagtataa aagaagactg 300
ctcctgggtca aagaanacag aagaaatgtg ccgagaggtc cttggaccag acgatatctg 360
aacaatacag aattgtcacc aaatgaaca 389

<210> 701
<211> 346
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 701

aacacaccaa aagattatga tgatgcatgg ctcaaattct caccactttc aaattgagct 60
ttcaaaaacta tcatgacatg tagaagaaaa ataaggattt caaatcagaa aatgtcaaga 120
gacttttatt ttcagaacaa ttaccattt cttgaacata tcctataatt caaagaanaa 180
tatgcaaagt tgtacatgca aacagaattt gacctaaata ttaaactaga aaccaacat 240
acttaaaaac aaaactaaca aaactaaca aactaggaat accaaactaa cttaaaaaat 300
tactaaacca aaaccaaaga acaagtcccc cataacttaa caacac 346

<210> 702
<211> 416
<212> DNA
<213> Glycine max

<400> 702

agcttggcat caaagcgctc tattcagcat ttgcactcca cccttctcgg ggatcttagc 60
agggtttttt taccaatacc ttctctctcc aatatttgct tgaaatacat tactttaaga 120
attaatttct aaattacacc attttctata gcaaagtcgg gttctgacat cccaacttga 180
tattttttta tcattctaca cattctctct ccattgtttt cttacaatac actactttat 240
atattaattc tcacactaga tcactttcaa caaattcacg aagatcgggg tcggacttcc 300
cttttttatt tataaaacac tctatatatt attaaaatta aatattatat tatataaaat 360
tatttctaat taatatagaa tttagctatc tattaaatta atttatggaa tattat 416

<210> 703

<211> 434

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 703

tgataagtnt atgggctaaa cttggtttga ttaggggtga ttcataatta gtcatcaaac 60
aattaacttg aanagttaaa tgacttataa gagtaatttg atgaaaatac tttttttgag 120
ccacaagatt atacgaaaga tttgataata ttaaattcga gcagaatatg cttcctcgag 180
ataaaaataa agaagcaa at gacttgagac aaattgcttt tgggtataaa atgaatgagc 240
agaattttga atctcttata gcggtaaaaa aatttagcag atgagttcan aattttta at 300
acaaatactt taacacctct tgattgggaa aaaaaaatct ttgggttgatt atctctaana 360
tccaagctcg taagtttata aaaaaagata aatatagagc attgaactat gctgatttag 420
gagatgacgt atat 434

<210> 704

<211> 420

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 704

gcttcttctt ggttctctcc ccattngaaa ccatcatttt tcttgagcac ttcattgaga 60
ggtgctgcca atgtgctaaa atccttcaaa aatcgtctat aaaaacttgc taagccatga 120

aaactcctca cctcggacac agaacttaggt gtaggccatt cttgaatagc cctaacccttc 180
 tgctgatcaa cttgcactcc ttttgaactc acaacaaaac caagaaacac aacatgggta 240
 gtacaaaaga tgcatttttc aagattggca tacaattggt cttttctaag cacagtcaag 300
 acagatttta aatgatcaat atgcaaata agtgaagtgc tatagataag aatatcatca 360
 aagtacacca caactacact ttctatgaac tctctcaaga tatgggttcac taatctcatg 420

<210> 705
 <211> 401
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 705

ntataagtgc gggctctggga gactaatgtc aagtgttcgc gatatgtgaa gatgatgttc 60
 caagtacttc ggatttggtc cgaccatgcc ctctgattt ccagctggga aattggcgag 120
 tggaggaacg ccccggcatt tacgcaacaa gcataatgta aacctttacg gttttaaaag 180
 ctctatagtt gggcctaggc ttcagagttt tcattttggt aaggctttgt gtcttttggt 240
 ttttaattta taatacaagg atctntcttc atctgttcct ggtctctacc cattctcatt 300
 catttgcacg tttacttctt tntctaaaac ggcagattcg atgacgagtt ccccgaaagta 360
 ctaatacctg ggaccctgtc atcaacttcg agcaagaaat g 401

<210> 706
 <211> 421
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 706

ctctacant tcatctctag catgcattnt tctttcttcc tttaccact cctcacgttt 60
 ggttttttag ggaaaaacat cataactaaa cgcgccacaa ggcaccccta tcgcaccaga 120
 tccaaatcta taacgatggg tgatcaagag gagacacagg aacagatgac agccgacatg 180
 tcggctctga aagaacaaat ggcctacatg atggaggcca tgtaggtat gaggcagctc 240
 atggagaaga acgcccgcac cgctgccgct gtcagttcgg ctgccaaaagc agacccaact 300
 ctcttggaac tgtgcccac ctccttaagc gtgtaggacg ggaagggacc actgggcacg 360

atggcaccct taccttgata caaccgagcg gctaccctta ggatngccgc caactactca 420
c 421

<210> 707
<211> 450
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 707

tcgtgatttt ctggcagcat tgtgataaaa gtccttttag acaaagctct cataacaatt 60
tcagggtgcca gtaagttaga aatgcatgac ttgatacagg aaatggattg ngaaattggt 120
catcaagaat ctatcataga ccttggacga caaagtcaat attggaaatt cgagtaagtg 180
caagatgtat tgaaatataa catggtaaaa tggatattca actgctttta ttngataata 240
cctgtgcttc ttagaatgta tctagtagta tagacctgat cattattatt ntaatttttg 300
gttgtcatgg aactgatatt cgtgaaggca taactctaga tttgagtaaa ttaaccaggg 360
atctatattt gagctccaat tccttggcaa aattgtctaa catgagattt ctanaatcca 420
tgatttgtgc tacatgactt actgatttac 450

<210> 708
<211> 330
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 708

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tcgtgtgatt ccccgagatg aaacaaacat tcctcacttg tgccttcgcc acgagagacc 120
atgcatgttg cctgcagcga ctggtagatg aaccgtctag acatagccac atcttctaata 180
ctctctgact ctgacgacct attctttgtg atggcattta tgctaactcc cccatgactg 240
gctagtggat tggtttaacg ttggggccct cttcttgaaa ggatagtctc tccacactta 300
ttatgtgtag caccttatac ttgaatggcc 330

<210> 709
<211> 381
<212> DNA

<213> Glycine max
 <223> unsure at all n locations
 <400> 709

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 gctcatgcaa caattgtag ccatggctat acgagacatc ttgccaaaca aagttagggt 120
 agcgataact cgcattgtgct ntttcttcca tgctatatgt agcaaagtca ttgatectat 180
 caagtttgat gagttggaaa atgacgccgc aattatactg tgccagttgg agatgtattt 240
 tccccctgct ntatttgaca tcatgattca cttgattgtg atctggacag agaaatcaaa 300
 tgttgtggtc ctatttatct accgaggatg taccgggtg agcgatacat gaagatcgta 360
 aaagggtata cgaagaatct a 381

<210> 710
 <211> 464
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 710

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 aaaagagagg acttgaacag aggtaaggta taaaacacca gttctgtggt gaaagtaaaa 120
 acgaaatatac tggctcacac ttgattcaat gaattgaata gctcaaggaa atataagcac 180
 ccatgatggt tgtggcatgt acttcaacaa ttaattcaat atagaaacta taataattga 240
 tacgaaataa aatgtgtgga aatattgaga ccatacttca natgagtaag taatatgcat 300
 cttgagggtc caagtatata ttgatggctt cacagattca tcccttgaag ttaaaatact 360
 aacaagcaat ttgaagatca actttccagg tatgtaaaac acagaattag gatcatctac 420
 agtcttatat agaactgcat caacctagtc cattgattta atta 464

<210> 711
 <211> 502
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 711

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tgataactca	gccttgtgat	cactgtatca	ccagcatatc	tttcagctct	attgaaagcc	120
tatctgcttc	aaagaaaaat	cgactcatag	tcagggcggt	taacaaaaat	cttctggaaa	180
ggactcattt	ttgaattcat	ctttaagggtg	ttcagtacat	gtatggngtg	agacaggttt	240
ggaaaataca	attgtttcag	accccaagcg	agttaattac	tgctcacagg	ggggtagagt	300
tatgctaaat	gtgtcagcag	atgggtacccc	acgcaatgca	natataatgt	ccactatttc	360
cgatgaatac	cagaagctga	agtactctgt	ctctcttgaa	atattttcaat	ttactttgtg	420
tgtgaacaaa	gagaaacagt	ccacacagat	ggaacttgaa	agaccagatc	tgtctatacg	480
aatatatgag	gagaaangcc	cg				502

gcttttttggga	gtagaaacat	gggaccaact	cattttattt	caaaaatgtt	gtatctagtc	60
aaggtctaag	attccataca	agtttcttag	cgatttctaa	ttatgtgggc	cattaagtct	120
atcatatgct	gacaatagcc	gagaaaccca	tgaatttctt	cgggggcgga	gtaagtgtct	180
gccatcgct	tggccttggc	taacaatcgg	ggaagttcct	gactcccggt	caaggtaaga	240
gcaaaccgat	ccatccacat	ggttgcctct	tggtgaaaga	gtcgatcacc	cttctcttag	300
cctctttntc	cgcgataact	tgggcatact	cgtccgcgat	cctatgctcg	tgggccgtgg	360
ctagacctaa	ctcttcttgg	tacttggcga	tgatagctag	catgttggtc	tcggtctcgc	420
atagacgctg	agacaagctt	cttttggacc	ttgaacaggc	aactaactcc	tctt	474

gtgcgggtct gggagacgaa ggtcaagtgg tcgcatatg tgaagatgat gttccaagaa 60
ctctggattt ggtccgacca tgcctcctg atttcagct gggaaattgg cgagtggagg 120

aacgccccgg catttacgca acgagcataa tgtaaaccctt tacgggtttta aaagctctat 180
 agttggggcct aggcttttaga gttttcattn tgtaagggtt ttgtgtcttt tgaatttata 240
 atacaaggat ctttcttcat ctgttcctag tctctaccca ttctcattca tttgcatgtt 300
 tacttctttt ttctaaaacg gcagattcga tgacgagtc cccgaaggta ctaatacctg 360
 ngaccctgtc atcaacttcg agcaagaaat gaaccanacg gaagatgaag gagatgagga 420
 tgtggga 427

<210> 714
 <211> 289
 <212> DNA
 <213> Glycine max

<400> 714

atttccccct tcttcttttt aaatatacca ttataaaaaa gggaaaaact tacgtagagt 60
 tattaaccta ttgtggaga tttaaacaat ctgggttaag agatccatct aagagcctaa 120
 ataaaatctg gaagacagaa caagggaaat gttaaaagaa gaagaagtga atttgcata 180
 atagaacgcg tgagagaaga aaaagaaggg acctcaattt gccattggc tgcagcaata 240
 tgaagaggag agtgacaatc atagagaagg gtcagagttc aagagagct 289

<210> 715
 <211> 383
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 715

tcttctgacc cccgcgacat attctcaagt accactcagt ggtcaactaa taaaacgtgg 60
 aggactgact ctttcacact ttctcacacc gagcttattg gggttatgggg caccggtcat 120
 atgtggtact aggtggcgat cgggcgatgg cgcanaacaa atatccatt tccacaagcc 180
 caggcataag cccaccatcc ccangtgccc acctttaaaa ttagttcatc accgggtccc 240
 catcaacctc tccaagcttt cacaatatct aaacaattca attcatttg tcatgaaact 300
 accttaaaaa cagagtgaag gtagaaatct ttacacaaga ttcatcana ctccacatag 360
 tttttccaac ccacatacct cag 383

<210> 716
 <211> 69
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 716

agcttgaagg tgtgtâaccc accattttcc atattatata tactggnaac gtgtctacta 60
 tcatggaca 69

<210> 717
 <211> 240
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 717

agctntgtat aatgagaatt gtccttcaac aagctttgaa anttggcatc tgaagtctga 60
 aagctctang cagataagtc tgcaaaagct ggaagtgggtg ctgaagtaaa agatgcaagg 120
 atgccagcta ttggtgcana ggaagaggga gcatcagctg ctctgatctt ggtcttcctt 180
 gcctctagaa aattaactgg ttggtcattc gcattccaac angttcttat gatataagct 240

<210> 718
 <211> 461
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 718

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 atctcatagg gatgaatgac tcgggcatac tttaagctta tgcacggaaa atgtaattat 120
 gaaattgaga tgcccgaaga aacaccattt cctagttaac catgcattag gtaccatggt 180
 caattatattt gttttgttgt tgtgtgtttt ttttttagaa atgggtttat gatcccaaca 240
 tggttgggtc atgggtgccta acacatgcaa ctaagaatgt agtgtgaagt ttcaogcttc 300
 ccttttttgg tttttgtttt gtagaggaaa acgcaaggat gagcaaacaat gaaaacaaat 360
 ggtatgcaat tntgcagatc aaaaagtttg ttgaacgcat atgcatgatg atgccatgac 420
 tcatgcanaa tgtgaggctg gaatatgata acggacaaat g 461

<210> 719
 <211> 398
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 719

ttgagggaga atgtagata atatgtttt tctttaataa ctttcttaat atacatcgta 60
 gactatatta acacattcta agtaatcgat tacgattttt ttttattttt tattttatga 120
 tgaatttcat gctgatcaca cacacttttt cactcatata atgagactaa aagaataaaaa 180
 tatacgtata cacgcaatat aaataatgga aaggaatata aatttactgt gagtcgacac 240
 tttcaattat ttttatgaaa tatatcaata aatattcatt atcctcaatc aattatgaag 300
 ttttagacg aattttctcc ttttctttgc gagacttctt tcatgtcggg cgatggctct 360
 ctttcaaga tgattattct tatttcanaa acttttat 398

<210> 720
 <211> 516
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 720

ccggtnatat ntttgaaacc ctggagtttg gctttaccat cgatnaccca cannanaaan 60
 aanaacacat ntttgatgaa acacgatgac acaatcacia cgacttgtgt catgtattgt 120
 actacggaga tccttaacga gtgaatctct ctctaggaga cacgtatgga actgtatgaa 180
 tattacgcat acctacttta gagcgaagct ctactgcttg agaagacaaa ctagatattc 240
 acacaacccc atcttgatat gtttagctca cccgatgaca tattacatgt taagtgtgct 300
 gaacttctgt tgtgctgaga cacatttatt gactctctat gataccgtcc tgcgctatgc 360
 acaggttata gagacttata ggtgacctgt ctaacgaaat gccagttgat atcgtgtcat 420
 ctataatcgg gcttatgatt ggacacgggc tcgaaactaa cgtatggcga agacaacatg 480
 tggactatct tgggtttatc tcaactgatct gggaan 516

<210> 721
 <211> 391

<212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 721

atcttaagtc acctgccgca tgcaagcttg cctcanagag gtccaggaag gacatggcat 60
 ccgaaggaac tagttccgct ccggagtatg acagtcaccg ctttaggagc gctgtacacc 120
 agcagcgctt cgaggccatc aagggatggt cgttttctcca ggagcgacgc gtccagctca 180
 gggacgacga gtatactgat ttgcacgagg aaatagggcg ccagcgggtg gcatactgt 240
 gtactcccat ggccaagttt gatccagaaa tagtccttga gttttatgcc aatgcttggc 300
 caacagagga gggcgtgctg gacatgagat cctgngtaag gggtcagtgg atcccgtttg 360
 atgccgacgc tatcggccaa ctctaagat a 391

<210> 722
 <211> 289
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 722

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 cccatagagc tccgcgaaga ttattccggc catactcttc cttgcgagcc ctcttggact 120
 cttggtcaag ggctcttgcg gtaattgcat tctcttcccg taacccggca cactccttcc 180
 gaatgtgtgt agcggccaac ttgaacttct ccttggaag ntgcgcttt cctaactcgc 240
 ttttgagagc tcggacttct tcgacctctt ccgaggcttc aaactctct 289

<210> 723
 <211> 296
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 723

atgcaaaagg tacgtatagc aacgtgcctt ctattcggtc aagcagactt gcacttcaac 60
 ttatcatatt tttttcttct taaacaacat ttttatccta aaataaaata aggactctgt 120
 aattntatnt tattttaaaa aaatattttt aaatctcata tattttaaag agatgatgaa 180

gtcagagtaa ttgattatgg aattttaaaaa tccagtttat aaaaaaaatg tgattatccc 240
atcgaatttc taaaaaaaaa catgacacat cagtgggtta ttaatattat tgggat 296

<210> 724
<211> 125
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 724

gtatgtgntt ggtatttatg ttaaataata agnttaanna taanaatgag agtttgtatt 60
aatatttaat agtatgtaag gngatgaaaa aaataaatat ataaataaaa tgaattnatt 120
attag 125

<210> 725
<211> 392
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 725

agcttcttga ggaagcctct taatgaagct tcttgaggaa gctacatgag ctgccttggc 60
aaaaacgttg cccagccttc gataaccgtt ggatcttcgt gtaatttggc ttgcagcttc 120
acaagacaat tgtacacgat ctgcctgttg ggatcttga gaagatgtct ggagtgtgtg 180
tgaagcttcc gttcccgaga gaatttctca tttaagcatt tcagcctttg ctttcgtgta 240
gcttaagaat tccttctcct ttctttcttc canagtcatt tctaacgccc caagcatttt 300
ctccatcacc cacaaccacc attagccatc acanaccgcc attgttctcc attgagaccc 360
acattgaaag gaacccttca accgaagcgg aa 392

<210> 726
<211> 447
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 726

ggaaggtttg tacatgacca aatctttagt taatcgtctt tacctanaac agtctttgta 60
tttgtttaag atgcatgaag atagatcact aggagaacaa ttggatttgg ttaataaaact 120

gattctagat cttgaaaata tcatgtcac tatatgatga tgaggatcaa gctttgttat 180
 tgttggtgctc tttgcctaag gggtactcta atttcaaaga gactntattg tttggaagag 240
 actttgtttc tcttgatgaa gtgcaggctg ctctgaattc aaaggaattg aatgaaagaa 300
 aggaaaataa gtcctttaca agtgggtgaag ggctgacagc aagaggcaag accttcatga 360
 caaatagtaa atctgataag aagaagcana agccagaaaa ccagaagaat ggtgaaggaa 420
 atgtcttcan aatcagaggt catcact 447

<210> 727
 <211> 329
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 727

gatgggtgaga ctgactctaa atttgatttc tgtggaccag tcttagtgct tttgaaattn 60
 tatcaaaata taccattgg ttttttatta aattgcgctt gaattaaact agataaattt 120
 tgcaatgaat tctctattag agccccctgt gagatgaggt caaattaagg tccatttggtg 180
 acatacaagg tggctgccct attgtgcaaa ttgggttttg cagtgtgtgg gtctctttga 240
 ttttgagtct ttataagggt cgaaatgttg tttatagctt cttaaattag ctcataaatt 300
 ctttctaacc atgtctatta gaaaaatta 329

<210> 728
 <211> 429
 <212> DNA
 <213> Glycine max
 <400> 728

agcttggtgg agttcaaaga gaatcgtaga aggttcgggc taagatataa gcctacacgc 60
 accgacatga agagaaacac cctataaagg agaggcagaa gtgtgggcca ccagcaagga 120
 ttgcaagtaa aaggaaactcc cttatgtcac atcaacaaga gttttgtcag cgcacgctgg 180
 atgtgtgagg ggtgggttgc catgatccat gatgaagtcc ctcaagagca atcaaactgg 240
 gtgcgggcat gccctcctat gttcgagttg ggaaattggc aaattatcaa acaaccaca 300
 atttttgtgg caaacataat gtaatttggg aatccaaacc ctatagctga gcctcggctt 360

gtcttttgc attagatata tataaaatat aatctggcctt ctttttttct tgcactttca 420
tccctattt 429

<210> 729
<211> 265
<212> DNA
<213> Glycine max

<400> 729

tctttgagaa aacttccttg agaagctaga gcttagctac acttaccct ctcataacta 60
agctcacctc cttgagaagc ttccttaaga agattcgtaa agaagctaga gcttagctac 120
acatacctct ctaatagcta agctcacctc cttgagatga gaagctagag cttagctaca 180
cacccttat aatagctaag ctcaccccca tgacaaaaaa catgaaaata aaaaaaagt 240
ccttattaca aagacaactc aaaat 265

<210> 730
<211> 254
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 730

ggcacctcac ataacctgtt attctctngn ccaagggaaa gatttggagt aaaactttgg 60
aggtccacga aacagacgac aaattatttc taatgtataa taaatacttt tagtttgata 120
tattttttta aatgaacgag aaaaaaatga tagattaaca taaatggaat gttctaacac 180
cccagtgcc ggaggcttcc cgctatacga aggtatgtgg gaggggtatt ngacacagac 240
ttacccttgc ctat 254

<210> 731
<211> 559
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 731

ctcgtaactt aactcggcctt cgtactcgca actgngctcg tatttctatc ctatatgtta 60
tcncacacgc tgtgtatttt gttgattctg ttcgaaccg ngatcctata natagacctc 120

acgcatgcc a gcttaaatac caccagcatc aaagatctat ggtctgttga cggaacctct 180
 ccaaatacgca tctttccgca agaattacgg aaagatctta taattgacct tagccgacgt 240
 atccatgaag ccattgcgac actcaaccta ttatacgacc agcctttgaa atgttttcat 300
 tcggagactc tcatctaccc caccatgga aatttgagaa acttttaggtg cctctccggg 360
 gaaaaaacat attttctttc cggggctccc tctttgacca attgccctgt ggcaagggtt 420
 agaaaagggtt gaagattaaa acactcgga cggctggtgc cccccgggg actctaaaac 480
 tggaagggtt ggcaccaagg atgggcccc ttttggggga ccctctattt ttggggggcc 540
 ccttcacccc cagcgtgc 559

<210> 732
 <211> 329
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 732

gagaaaactt ncttgagaag cttctttgag aaaactttct tgagaagcta gagtttagct 60
 acacacaccc ctctaataac taagctcacc tccttgagaa gcttccttga gaagattcct 120
 aaagaagcta gagcttagct acacacaccc cctataatag ctaagctcac ccccatgcaa 180
 aaatacatga aaatataaaa aaaaagtccc tattacaaag actactcaaa atgcccttga 240
 atacaaggct taaaccctat actactagaa tgggccacat acaaggcca aaagaaggaa 300
 aaccaattcc tacatttacc aagaagaat 329

<210> 733
 <211> 470
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 733

acactcttag gatttgccta gtttacatth cttgcttaca ttcataatag cttatttcct 60
 ttaccttcca ttgtcaaacc gcctagatag ctttcctttt accaattagt tntttacctt 120
 atctttcaca cctcttttag tgtttatttg gctagnttca accatagttt cttttacctt 180
 ttgttntcaa acctccaaca agaaagaacc acaacttagg aaccaatatg agtcatcatt 240

catctagtg taatggcaag ggtactagtc ataaagaccc tttatctaga atcttagatg 300
 agttgagttc cctcacgtta tggaaagaan aacaagagag aaaagaanaa ggaagaataa 360
 gagtggaaga aataaatcat gatgaaagaa agacaatatg agaggaagaa agaagaacaa 420
 taatgaaaga aatgaanaga gaaaaacatg cctnctatag tagtcataac 470

<210> 734
 <211> 429
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 734

atgatttggga ttatcctcta nggcaatcag tattcagtat tttaaattcc ctctcaaaaa 60
 attgcaggct tgggtggcggg gccacaaagc tgccatatat tataagaggc ttagtagagg 120
 tgcaatagtt acacaatgca gatggagggg gcgcatagcc aggaaagaac ttatgaaact 180
 gaaaatggta tgttttacca tgattcttat acattaagta gatccttcag aaatagatga 240
 caaaagatgt aacacgtcca tcttcaaaaa cttatggcac aaaaagcaaa agatactatt 300
 tgatcaattt ttaattgtaa aattgagttt caaattatag atatatacaa acaatattca 360
 tattttttgt ttcattatta tcttcatgaa gggaaaaaca atagaagaac ttanaattct 420
 cttctgatc 429

<210> 735
 <211> 371
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 735

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 atctttcggt gtaactccat ggctgaanat caccattgaa ggacctcatt aaagctcaaa 120
 gatccaacct ttataaaagc ttctcaagaa agcttccatc aggacttttg gacatcatag 180
 tgcctacagc tgaggcatca aggatcatct tgggtngagg tttcatacca ctatggaaga 240
 tatgcatatg tgcactgtca tcanagttat ggtttatgca cctctgcaac aaagctttaa 300
 atctctccca tgtctcacag agagggttgt gagatccttg ggataaagtc atgatgaagg 360

ggtttacact g

371

<210> 736
<211> 335
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 736

gagaaaagtgt ggaagagtca gtcttcctac ttttattcgt tgaccacaga gtggtacctg 60
gagatatgtc gcgngnggtca agagaccttg tggacgtcaa gtgggggtgtt attgccc aaa 120
accaagcttg accaatcccc acccaacca ggcatagtca gtcagtgaga acctgtgacg 180
tacctaaaca ggcgagctcc tggcagtcaa ccgataaaaag aacaaagacc acaaagcaag 240
gaggcttgtg tgggtggtgg ccagctatgg atcttgagtg atatatgggt tatggcctct 300
ggtaatcgat tacaaagggg gtgtaatcga ttaca 335

<210> 737
<211> 504
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 737

atccttaagt cacctgcggc atgcaagctn gttcgcacat cgtcgcgtg tatgacatcc 60
actccacaag gtttgaagta gaggagacct tcaatcctat tacgcaacgt ggcggacaaa 120
aatgggcagt taacttgaat ggtcattatt gtcaatgcgg aaggtattct gcgcttact 180
atccatgttc acatattatt gcagcttgtg gttacgtgag cctgaactac taccaatata 240
tagatgttgt ttatacaaat gagcacatct tanaagctta ctccccacaa tgggtggcctc 300
ttgggaatga agcggctatt cctccttcta atgacgcatg gacacttatc catgacccaa 360
ctacaattcg tgcgataggt ccggcaaaat caacaaggat aatgaatgag atggattgga 420
tcgaaccatc tgaccaccga ctaaaatgca gtagatgtgg agccgaaggg cataacangc 480
gtcgtgccc atgcaatctg agcg 504

<210> 738
<211> 411
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 738

ttcacgcgacg aaaggaccac agtaggtcta ataagagaca aatctgatca tcatgctttg 60
atacatgccca aaaaaaacta cggcaaataga agagggtgag aatgaggaag aagcccatgc 120
tgtgactgtc attcctatac agccaagttt cccaccaacc caacaatgtc attacttagc 180
caataacaaa ccttttcctt acccaccgcc agttatccac aaaggccatc cttaaataca 240
ccataaagtc tgtctaccgc acttcanatg acgaacacca ccttttagcac ataccataaa 300
cacciaacaa gaaatggaat ttgcagcgag aaagcctata gaattcacc caattccagt 360
gtcctatgct gactngctcc catatctact tgataattca atggtagcca t 411

<210> 739

<211> 420

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 739

agctctcatc tagccaagaa ttacaaaag tgttacaaca taacctaacg atttctaatt 60
atatgggcca ttaaatttat catgtgttga cagtaattga ttagcccgtag aatttcctct 120
ggggctgaac acacttcggc catgggcctt gctttggcta gtagtcgagg gaggtcttga 180
cttcatttta aggtcaaggc gaacctatcc atccacatgg tcgcttcttg atgcaatgca 240
tcaatcacc tcctcttgc ttcttctcg gcgtatgctt gtgcgaagtc ctcttctatc 300
ttttgctcat gggcanaga ctgggttaac tcttcttctg actgtcctat tatanctagc 360
atgctctgct ccgtggcttc taagtgttgg gccaaacttt tcttggatct tgagcaagct 420

<210> 740

<211> 427

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 740

tgtatcgatt cgctcctagt caggatcaca tcatcttgct cctttctttn tcatttaact 60
tttggctttt gtttccttct ctctttatat atgttttggg gtataataat tcatatatgt 120

atgtatgtgt cggaacctac ccttcggtaa gagggcgagg cgaaaagcca aaggagcatc 180
 ttccaaaaag gaaaacccgc gggagtcgcc accaacgttt actctaggaa aacattagaa 240
 aaacccaaaa aaaaagggtcg aagggtctgca aattttgaaa atgaggggttt gggagttggt 300
 tacacacgag gaaggtattt gcacccacg cactcgtcac aagggatggc aacctttaat 360
 cgagtgtgca naacatgaac ttcaaatgtgtattttccc tttcatatnn gtttttttat 420
 ttcttttg 427

<210> 741
 <211> 500
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 741

ttaagtcacc tgccgctgca agcttatgat ttttanaaa ataattactt ctctgtgagt 60
 ctccatgaac ctaataaata tgaggagaat gaagatttgt gagtctccat gaacctaaaa 120
 aatctgagga gaatgaagat tggggaatga attataaaat tgtgatcctg tcttttataa 180
 acctaaactg acccgctntt ttcaggaaaa aagataagaa tagttaaact ttgcccgtat 240
 catacatatc actctaatan aaaataaagc attggagaga ccagaacata tgagatatga 300
 atcgaattct tatgcgttta tactttacca taaaagtcac attcaatatt atgtgaacca 360
 ttgatntaga tggctttntt cttataaact catcatttaa cagagttcag aataatataa 420
 attgatgaat aaatcacatt cgcgcttctt atatactgaa cctattaatt caagccttaa 480
 gctaataataa gcaacatttc 500

<210> 742
 <211> 412
 <212> DNA
 <213> Glycine max
 <400> 742

aatgtaataa gttaaacata aagctattga ttgccgtaac caattggcct aaaggccaag 60
 tagagtatat taaatttcag gcaaaccctt atttcgaatc aactggaagt cgatcagcaa 120
 atctttttaa ttaacgaatt tgcctacttt caggaaggaa aaaaaaaaaa agcagtgtaa 180

tacctaatat atatctgaat ttatcatacg tatttgctac aattagattt tgtaaactgt 240
 aaatctctct aatataattc ctttatgaaa gatgattgta acattaaaaa atatattttg 300
 ggtcagcagc ttggatcttt ataggggggtg gtgccaacct agttgggact actccctcga 360
 ttgtatcagc tgtctcacca ttcatttaat aaataaataa ccaggatcaa tt 412

<210> 743
 <211> 361
 <212> DNA
 <213> Glycine max

<400> 743

gagactatgt ttggcaaaga aaaactatga agaatgcgtt gccatggaaa aactacaacg 60
 cctactctct tcaaaagaaa taaagaaatt tcaaaaattg aaaaagata tgctgaagag 120
 gtgaatgaat taaatgataa aatccaagag atagaagata aatattatgc aacagagaaa 180
 tggtagcagc agaggaaaaa cactattaag aaattattgg agtacaagtg tgaggtgaag 240
 tcccacatta aatagaagtg gaaaagttga gcaccatata agtgaggaga agacctataa 300
 atctaagtct taaggttttg agttaaagtg tgggtattaa atcccttatc ttgttactca 360
 t 361

<210> 744
 <211> 386
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 744

tggtcaccat tntcaaggat agaattcata agactaagga aaaggattaa aaggaggtca 60
 atgatgatga tgatgatgat gatgaagact atgtacctaa atatgaagag agacttggat 120
 caaattctat aagtgaaaat taccagcagc acaaatatga tgaattttct actacaaatg 180
 atcttgagtc acgggaaaga aagataactc ttgaaaattt tcaatataat gctaataatac 240
 attattgtgt taataaatga gtgggggtggt tttgtgggta ttgtgaaaat tntcaatata 300
 ataccatagg aaattgactt cattntcttg atctatctac atctttctta anaaaaattc 360
 aaattcactc ttttgtaana aaatag 386

<210> 745
 <211> 244
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 745

cataaaaagt cttgtggaat tgattacaag gattnggtaa tcgactacca atgacaagtt 60
 ttgaataaaa atcacaagat gtaactcttc aaatgggtttt caggctattc tanagggtat 120
 aactcttcca atgggtttcca ttgaccacac ataaagagtc tataaaagcg cgaccttgag 180
 tngcatattg agatctgagt acaaactttt acatctttta cacacaacct ttgaacatct 240
 tctt 244

<210> 746
 <211> 451
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 746

ggcttctaca atctccnct ntttgatgat gacaaccctg atatcaagaa acacatgcac 60
 atactttttc ctagtgcatt actcaacttaa ttctccatat tctaccctt tgggtttgag 120
 tttaagcttc acttgaaatt aagttaatta cttatgtgag ttcttgattt aatccctatt 180
 tctctcccc tttggcatca acaaaaaacc aaagtgcgta acaaatataa aacatacaca 240
 aataactaat catacacaag acattcattg aaaaatctaa accaatcatg aagcaagaaa 300
 catgaataga tcaaatatat aaaaaccaca tagtcatata acataattca tatttgttca 360
 gtcatactat gcaaataaaa gaaatactaa atgttcaa atgtcataataa tatagccaaa 420
 tacacggcta gaaatcaaag tactaataat a 451

<210> 747
 <211> 476
 <212> DNA
 <213> Glycine max

<400> 747

cgcgcgcgcg ctcttattag agaacggtaa cgtcttgtag tattatacac acgaaactat 60
 atctacagga tgccacgtat gagggtattac ctcaataatc aggcgcttag actcgatctg 120

agagaggaga gacgtataaa ctcatcaact tctctaggat tgaagttctg atctatatcg 180
gaacattatc ttctgacgcg atacctgcat acactacgct cagcatacca gacctctctg 240
cattgtatac gacggagggtc ctccgagtag agttatcaaa cataacctat ccaattgaag 300
cacatggact tttcaataaa aagcacgaca attccattat tacacgtctg ctcatcgaga 360
aagagatctc taggccgtag atatatactc tcatattcaa cttagatgaa tgtagaaaaa 420
atgtacttat agctgcataa tgatcatgtg ccgcatctaa caatacacga gatgcc 476

<210> 748
<211> 337
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 748

tctccaccat tntcttataa atagggggag aagtgaagag gaatttcgtt cagccctcct 60
ggtaattcag aatcacttaa aattagtgaa naaaattggt tccgtgaaga anatccaagc 120
cgaggcgctt ccgtaacgtt tccgtgggtg atttcgcaa ggttttcggc cgttcttcga 180
cgctcttcat tegtcttcg tcgntcttcg gtcttcaacc ggtaagttcc ctaaactcaa 240
cttttcaatt cattctatgt acccttagtg gtcttcattt gcttttacgt gctttcattt 300
acatttcctt tacttttcgt acccccgttt gacgtgc 337

<210> 749
<211> 489
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 749

agcttgtctc agcgtttagg cgagacagag accatcatgt tagctatcat cgccaagtac 60
gaagaagagt taggtctagc cacggcccac gagcatagaa tcgcggatga gtatgctcaa 120
gtatatgcgg aaaaagaggc tagaggaagg gtgatcgact ctttacacca agaggcaacc 180
atgtggatgg atcggtttgc tcttaccttg aacgggagtc aagaacttcc ccgattgtta 240
gccaaaggcca aggcgatggc agacacctac tccgccncg aagagattca tgggcttctc 300
agctattgtc agcatatgat agacttaatg gccacataa ttagaaatcg ttaggaaact 360

tgtatggtct cttatacctt gactagatat gaattccttc ttgaaatana atgagttggt 420
 cccatgtttc tactccacan agcttggtgca aatcanatca ctctacatc tcattcttag 480
 catgcattt 489

<210> 750
 <211> 451
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 750

ggcgaccagc tcgcccaggc gagtaagggt gcttcctcca gaagcaacaa ccttctggag 60
 gaatcttcta gagggcccaa gtgggcctga ttgctatttg tacctccctt ttactaaat 120
 ccatccctt ctattttttt ggtaattctt tttccgtaac gttacgaaac ttacgaatt 180
 tcgtaacgat acttattttc cttccgcaag gttacgaatc cttacggatt atgtatttac 240
 tctnttttag cntcgaaga agttacggaa acttacggat tgcgcanaaa cacctctttt 300
 cgatttccgc cacattacgg aatttcacgg attgcgcaag cctgcttctt tttgattntt 360
 gacaggcctc gggacttcat tcattgtgca accaaggacg ccaagtatct cgaagcggcc 420
 aatcaaagggt tgtatatcat caaataataa t 451

<210> 751
 <211> 141
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 751

tactgaccat tgataatatc acaagtgagt ttattcagaa attagagttt atgtctttat 60
 cttgtgagag tgattctcct aaattcttga gtgattanag aacaccctgc ctgtatcaaa 120
 ggactttaac aacctttgag a 141

<210> 752
 <211> 131
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations

<400> 752

gattacctct ataatcaggc tcttagactc gatctgagag acgagagacg nataaaactca 60
ttatcttctc aaggatagag tgtctgatat atatcccaac aatatcatct gaggtcatcc 120
ctgcatatac t 131

<210> 753

<211> 427

<212> DNA

<213> Glycine max

<400> 753

taggggcagg tacgaacagg ttctatgcc atgatcaatc gatcaccacc cccgcgttcg 60
gctaaagata ttaaagaagc tctcctagga ggcagcctag tatctctaac tttgctcttt 120
aatttctctgt ttcatacttg ttctttttct tgaactatat cctgaattcg cctaagttta 180
tatgcaatta taggatttta agagaaaaaa tataacaatg aataacacaa ttttgtaaag 240
gattttcttc accaaaaaaa taataattac ctgcgttggg cgagtggcca gctcgcctag 300
gcgagcatgg ctatggtgaa aacataaaa aggggagggg tgaagccatt ttcaccctat 360
tcttgcccaa aatcaaaacc ttccccaaga gcttacggga gccaccattg gcagcagccc 420
ccaagct 427

<210> 754

<211> 411

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 754

catcacatgt ggcactatgt ggcggtcggg cgatggtgca caacaagttt tccacatcca 60
caaatcgcg ataaaccac catcccctgg tgcccacctc caactgagct cacgtactcc 120
cacgtagccc atattctcgt ttctctcaac accgggtccc catcaatcct cccaagcttc 180
cccaacattc aggtaattca acatccaaat catcacaac taacaaacca agcaaaacag 240
ggcaaaggca gaatactctg cccaaaactc aaaccanaat cacagctttt tctcacttaa 300
agaccccagt aacatttcct tcattccaat tcgttaaccg gtggatcgac tcgaaaaatt 360
tactggaagt ctctagtaca taagcctaca ttntgaccgg tgggatctac t 411

<210> 755
 <211> 268
 <212> DNA
 <213> Glycine max

<400> 755

catgcctttg acggcaaccg cccacacgcg acgtgagaga tcgacctccc tgtacagata 60
 agcccccata cctgtcatga tacccttcaa gatattgata ttaacctggc ttacagctgt 120
 cttttgggac gcccggtgat gcactcagtg ggagatgac cctctacact gcaccacaaa 180
 ttgatagtcg tagtacacgt gcactggggc attgtgtctg tggacgaaga aatcttggtg 240
 agatgcccac tctctatgcc atatgtgg 268

<210> 756
 <211> 448
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 756

gcttgcataa aataatcaca gtatctgcat attcaattc agtatatgga catgtgttct 60
 tccttaagag aggggtgtat cctgttgaa gatcctgcat actacctgca ccaaaaggaa 120
 cataacacag tcattaagaa tcagtgttct gagtatatca atgaattntg tgtccacaag 180
 cctgtggcag aaccagtggc ttgtgtcaag ggaccaaggg attataattc ttcacgtatc 240
 cagtcgtgct taaaagttgt ctcaaagtga gatgaagagg ctgatgttga acctacatct 300
 cctgaanagg aaggataga atgtgataat ccagaatctg aaagtaggtg agaactctgac 360
 atcaagtgct gttatggttg tgaaactgat atataattgc atgacgtctt atagcagcag 420
 aatctgaagt gaaacttaat agcctaata 448

<210> 757
 <211> 454
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 757

agttagggta gtgaatgata ttctgaaggc gtgttattgt gtaaccatt aataaagtgt 60

ctcgattntg cttataaact ctgaaatgtg ctttgctctt ttgctttaag attntatattt 120
 attttggtac agaagatttt atttttttga aacaacaagt attgacaact ttctggctgt 180
 agtaaataga agaaggaaag tggaagaaat aggaaaatga taaactgcac ctctttttac 240
 ttaatcttaa tgattctcag ttgttttgct tttcctcttg caattgtctg aaaccaaggt 300
 ggaggtagaa gtttaaattc cataaagcac ttaaagaaac tatttaatat cctttgcgtc 360
 atttcatttg gatgggaaag ctatatacat ttagctagag catgccatta ctgcataaag 420
 aataccctaa ttaaggatag attatgagac acac 454

<210> 758
 <211> 88
 <212> DNA
 <213> Glycine max

<400> 758

tgtattgagc attccttttc tgtatcttcc gttttgctta gtttagccct gtaattctaa 60
 tatagattaa gagagcattc agcttgac 88

<210> 759
 <211> 172
 <212> DNA
 <213> Glycine max

<400> 759

tcttacatat taccctattt atgtctcaag atttagtgat tcaagcttgg cctcttggtt 60
 gagctcttaa catagtagag gaaagaaaac tcccgaatg atggaagaag gatatgctgt 120
 gaacaaaacc catatgttca agggagccac ctataactat gggaaggaaa aa 172

<210> 760
 <211> 315
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 760

gcttcttacc caggcaattc ttgggggnga agctccttct tccttggtt attccctagn 60
 ggatggtgcc tcccctatcc tcttctcctt tgccttcgc tgcctctcca tgatgaaaaa 120

tcaccattga aggacctcat tgaagatcaa agatccagcc tccatagaag ctccacaagc 180
aagcttccat caagttatga ccatttgaat ttctcgagat cttccgtggn tcaatttcgg 240
gcgtctccat atgtcatgtg cctgaatcgg acctccgtaa tataatttat gaccattcga 300
acttctctag agctt 315

<210> 761
<211> 410
<212> DNA
<213> Glycine max

<400> 761

gcagatctgg tcttcgccag tgaaaggatc aatgtgggtc cgaaaagagg caaatttgat 60
catcctacta tgacgactga gaaaactggg gcaaataaag aggggtgagga tgagggagaa 120
acccatgctg tgactgccat tctgtacgg ccaaatttcc caccaacca acaatatctt 180
tactcagcca ataacaaact ttctccttac ccaccacca gttatccaca aaggccatcc 240
ctaaatctac caciaagtct gtctaccgca cttccaatga cgaacaccac ctttagcaca 300
aaccaaaaac accaaccaag aagtgaattt tgcagcgaga aagcctgtag aattcacccc 360
aattccagtg tcctatgctg acttgtccca tatctacttg ataattcaat 410

<210> 762
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 762

gcttaataac aatgcctaag gaggagcatt acatagcacc catcttggtc ctgttaggga 60
taaacatgtc gttgaagtac attcaacatc tgggtataaa ggacttttac catggcatac 120
tacagtgact aacactgtat tcagaatcat tntcatccct tttatgaact tccctctttt 180
ttatgaactg cctctttttt tagtctttgt ttcccacctc ttaaacaatct gcattntccc 240
atthaggttt cttgtttttg tgaggaatgg cggaaattat gttttatatg gacaagactg 300
ttgggttgaa ttggaatgag aatcaacatg gtgctcatac ttttgttttc aaataaataa 360
aagtgggcat gctactaata tttaatntat tatggtgctt ctattaataa agtacacctt 420
gttaggaaac caaaatatag aacagaaat 449

<210> 763
 <211> 287
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 763

tcttatccaa cgctcatctt ggtggtgaag ctcttcttc catggcttat tccttaatgg 60
 atgggcgctc ctttcacctc ttttcctttg tcttcgcta catctccatg gtggaaaatc 120
 accattaaag gacccattg aagctcanag atccagctc catagaagcc ccacaagcaa 180
 gtttccatca gaatgtccac gtcttttagag ggctacacgc ccatgccttc agaggactac 240
 acgcctcgc cttatgagga ctacacatcc tcaccttag aggacta 287

<210> 764
 <211> 435
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 764

ctgtgagaac ctgtgacgta cctaaacagg cgagctcctg gcagtcaacc aataaaagaa 60
 caaagtccac gaagcaagga ggcttgtgtg gcggctggcc agctatgtat cttgggtggt 120
 atctggaaat tagcctctgg taatcgatta ccattcatgg ataatcgatt acaggggtta 180
 aaaatggaga caggatggta aatggcctct ggtaatcgat taccaaggga gtgtaatcga 240
 ttacacaggg tgatagggca ctggtaatcg attaccagct ggggtgtaatc cattacacag 300
 ggtgataggg cactagtaat cgattaccac ttatgtgtaa tcgattacac agtgtatattt 360
 ttaattttca atgtgcanag gctgtgtaat tcgtttttgg caccggtaat cattacatac 420
 tttggtatcg atacc 435

<210> 765
 <211> 318
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 765

taccaattga aattctgata ccaatgccag atgtcgtaca ggatgtcacg acatcacgct 360
tcagaacatg cagattatct ctgagtggat gaacacgata aacaagtata taacacaaga 420
gaattgttta cc 432

<210> 768
<211> 429
<212> DNA
<213> Glycine max

<400> 768

tgtctcagcg tctatgcgag acagagacca acatgttagc tatcatcgcc aagtaccaag 60
aagagttggg tctagccacg gccacgagc atagaatcgc ggatgagtat gcccaagtat 120
atgcggaaaa agaggctaga ggaaggggtga tcgactcttt acaccaagag gcaaccatgt 180
ggatggatcg gtttgctctt accttgaacg ggagtcaaga acttccccga ttgttggcca 240
aggccaaggc gatggcagac acctactccg cccccgaaga gattcatggg cttctcggct 300
attgtcagca tatgatagac ttaatggccc acataattag aaatcgttag gaaacttgta 360
tggtctctca gaccttgact agatatgatt tctttttttg aaatgaaatg agttgggtccc 420
aggtttcta 429

<210> 769
<211> 466
<212> DNA
<213> Glycine max
<223> unsure at all 'n locations
<400> 769

gtcacctgcg gcatgcaagc tngagaatat caatgcgtca nagtcgctat ctcaatcacc 60
tttgtttgaa gatggagttg tatcaactca caatggagat gggaggagat ctccatgacc 120
acatcaacaa gttcaatcgg ctagtaagtt aactgttgaa tgtggatgat aaattctcta 180
atgaggagca agcgtctctg ttgttgggtct cactaccaa gtcttcata gctttgggtc 240
aaacgttgct tgtgggaaga tcaactttga atttggatga ggtgactgtc gctcttagag 300
aanatgatga gaattgaaa tgctgatgat gaacacaatg caatagctgt gatggaatct 360
gagcgaggga ggaatcattc aaggagacat gatgggtctaa gaggaagatc acaatcgcaa 420
tcgcatccac aacgagatat gagtaacatt cactgcttct attgtg 466

<210> 770
 <211> 345
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 770

ctgaggatag agacttccta agctatatat cttctctctc anataagttc tctaactttc 60
 tagctatctc actctaagaa gtggattcac tcttgtcttg gatggttagg aatgaaggct 120
 cctaccctta tttatactac tccacctcca caatgaatgg tggagattac ttgtatccta 180
 ggggtggagat taattctcta gaattctcca cacattctag gagtctctac acttttctac 240
 tctctttcat atcattccat aaggtttcag aaggttccac acatctccaa aatatttcag 300
 agggttccac attcttccac aagcttctag agagttctac actac 345

<210> 771
 <211> 465
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 771

agctntgcaa ctcagacacc tcaacaggag agatccgacg aggctgttgc tcctcttgag 60
 cctacacctg cacaggttga accagtgcc a gctgatccac attctccagt ggcagatcca 120
 tcttctcccg aacttgaagc agctccccca tcttcaccta ttattatcat ctctgaagac 180
 cctacagagt caacatttgg agaagctggt gctctctctg attcccctgt tnttcatctg 240
 atgaatgagg aggagacaca ggatcagtca caggattctt anattcctgt ccttctgttt 300
 atgttgacaa ttatcataac tattatattn tagtacattg ttttagtgaa ttttggtgat 360
 gggttatatat acttgtgttt tttggggaaa gtacgatgca tggtttgaag catacaggaa 420
 tagttaactt gatcttgatg aatatgtagt gaaacacttt tatca 465

<210> 772
 <211> 397
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 772

tcattgcagt cttggaagcc agattagttc atggatgaga tccccacttt tgttgctcaa 60
cctttgcctg ttggaatgga tctggtttgg ttttggtttg gggttctcaa ttgaatttga 120
attcctctta attggttcct gcaaaattgt gtttttttgg gttcggttgt gtgttgatgat 180
aatgcaaaag gttgttaaaa atgattttta tttcattttt ttatgtgggg ggttgagttt 240
ctaggtgatt tgtatgaatg cacttgggtct tggagaaaaa aaattgtgtg aatttttagtt 300
ggaatggtcg aacagtgtga aaaagtttta ggttgcanat acacaactgt ggggtgtgatt 360
atgccaaaag gtatattaaa caaattgaat tttatgt 397

<210> 773

<211> 468

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 773

ttgtcgtgag aaaaagaata cactccaatt catcacatgg taaatgaata acaatggagg 60
acaaanagta tatagcaata ttaataaata aataaaaagt atatagccat tgaacgtagg 120
tgacaaattc aatccatgtc agtagtcgca tgacattcct ttttcttcta gaactgaacc 180
aacatcaccg aaaatcaaag ccattagccg aaagcanttt aacatagaaa cctataaaat 240
tgtgagttag ggttttagtga taggtcatgg tgagttgggg tcttggatcc ggttcaagta 300
tggaaccttc tttccatttt ttttatcttg actctcatga attaatgggc cagcaacact 360
tgatttgtgc ttgatacatt ataactaanaa ttagaaacct ataaaacctt ctcttttctt 420
tctcttttct ttttattatc tctttctaat ttttcttctt cctctctc 468

<210> 774

<211> 444

<212> DNA

<213> Glycine max

<400> 774

aaagatctca gttttctatt atataatgat ataaattagt ttaaaagtgc atcagatatc 60
agttttctat tttataatga tataaactaa aatacctgtt tttaacactg tcttcatggg 120
gtaaaggcac agtccatca tcaggttctt cgtcactctg taccattacc gcttgatcta 180

ctattcctaa gttgtgaagc cattctttta acttcacaag acagcggtat tacattagca 240
 attaggctga atcagctata tcaaatgtaa tgaaggaaac taaactaatg gagatatgta 300
 tcttcactta ataacaacgg agtaacatga taaagaatca gggtttggaa cctaagttga 360
 agtgtgaggt tggggaggcc aaaaagatta aatgatagac acaatactgc aaaattaatg 420
 tgcaaacctg ctcttgcttg tcat 444

<210> 775
 <211> 456
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 775

aacccatgga agctccta atctcccaca ctttgntggg tgggccattc ttggatggcc 60
 ttgattttct caaggtccac ttggaccca tttctaccaa ctacaaaacc taagaagact 120
 atattatcta cacaaaaggt acacttctct atatttgc atagagggtgtt tttcctaagg 180
 actgaaagaa cttgcctgag atgttctaag tgatcatcta ggctcctact gtacactaaa 240
 atatcatcaa aataaacaac tacaatcta cctatgaaat ccattaagac atgatgcata 300
 agcctcataa aggtgcttgg tgtgttagtg aagcccaaaa gcatcactat ccattcatac 360
 acaccatact tggcttgaa agcgcggtcc actcatcact ctttttcata ctggattcgt 420
 gataaccact ttaagatca tttttgaag agatat 456

<210> 776
 <211> 442
 <212> DNA
 <213> Glycine max
 <400> 776

gagctgaaca cacatacctc tataatagct aagcacacct ccttgagaag agaagctaga 60
 gcttatctac acaccccta taatagctaa gctcaccccc atgacaaaaa acatgaaaat 120
 aacagagaaa agtccttatt acaaagacaa ctcaacatgc cccgaagtac aaggctaaaa 180
 ccctatacta ctagaatggc caaaatacaa ggcctagacg aaggaataac ctattcta at 240
 atttaciaaag ataagcgggc tcatacttag cccatgggct cgaaatctac cctaaggctc 300

atgagaaccc taaggccttt ccttggatct ctagcccaat ctacttggag tcttctagcc 360
aatgcccttg cggggtaaga gtgcatcatt acttttact cagatgtgcg attcaggcac 420
atcagatatc gagacgctcg aa 442

<210> 777
<211> 286
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 777

gagcccgagt agtcaaagag aagttcaagt ctatagccat canagtctga agagagtatg 60
atgaactaag ggacgtcaat atggccaccg atgaagcctt ggaatgagaa accaagaatg 120
ccctgaagga agaacacgac caatacaagt tttgaggggc tttatagggc aataatagtg 180
agctcatact ccgaagaggt gaaaggagtc atcacgggtc acaggtatga tctgtaagga 240
cgagctatag gcttgcctta tgacgagaag aaatttgtcc cgaccg 286

<210> 778
<211> 193
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 778

atcagaccac ttccagngtg ctggaactac ttcacactga tttgatggng cctattctag 60
ttgaaagcct tggaggaaag aggtatgcct atgtcgatga ggatgatatc ttcagaatta 120
cctgagtcaa ctttatcaga gaaaaatcag acacctttga agctttaatt gagttgattc 180
ttatacttta aag 193

<210> 779
<211> 281
<212> DNA
<213> Glycine max

<400> 779

tcacacttac aaaggatata tgggtccatg agggacctcg ggctttctac agagggcttg 60
ttccatctct tcttggatg attccttatg cagggattga tctcactgca tatgacacct 120

tgaaagatct atccaagaga tatattcttt atgacagtgg tatggtatta ctgcaaccac 180
 attatctctt gaacttaatg gatttatctt accactctga aatttttagt gacacataac 240
 acatgtaaac tcaacctttg aacttaaata tgtaattttt t 281

<210> 780
 <211> 247
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 780

ttcgtacta aagttggcga cnttgaaaat atttggagta gtttttaaca gattttatat 60
 ggatattgtt gaattagtgt actatattca ttctatatat atgggcgtgt ctcaaataa 120
 gtgcatcaat aagtgttcca gaagggttac gagtaaagaa aagccacgt gattcccta 180
 actatttcgc gttcatctcc acgtgactcc cccaaggctg caactctact actaccatgt 240
 gctagaa 247

<210> 781
 <211> 118
 <212> DNA
 <213> Glycine max
 <400> 781

agcttggttg attatggcgc acccgtcata tgtggtacta ggtggcgatc gggatgatgg 60
 gcatatcaat tctttcacat ccacaaataa gacatgaacc caccatcccc agttgtcc 118

<210> 782
 <211> 260
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 782

tcagacaaaa gcaacatana atctatgtat ccgaaacccc tcaatctaata ggattgtcaa 60
 gggttgagaa gtgaatatga caatgagcgt tatttggagc aaactctcac ctacacaag 120
 tctataacat caatctaaac ttgctcaaac tggatttaca cctaaaactc caccgaata 180
 aaatttgatt cctcatcacc aattttaccc tagaaatgac tcttcgttca ctacgtacat 240

ccttttcttt tattgcaaag

260

<210> 783
<211> 449
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 783

agctntgcgg aattggcctt cgctggcgaa atgatcgaag tgggtctaaa aagaggaaaa 60
tctgatcatc atgctttgat aaatgcaaaa aaactggtgc aaatgaagag ggtgagaatg 120
aggagaaaac ccatgctgtg actgccattc ctatacagtc atgtttccca ccaaccaaac 180
aatgtcatta ctcagccaat aacaaacctt ctccttacc accacccagt tatccacaaa 240
ggccatccct aaatcaacca caaagcctgg ttaccgcact tccaatgaca aacaccacct 300
ttagcacaaa ccaaaacacc aaccaagaga tgaattttgc agcanaaaag catgtataat 360
tcacccaat tccggtgtcc tatgctgact tgctccata tctacttaat aagtcaatgg 420
tagccataac ctcaaccaag gttcatcaa 449

<210> 784
<211> 399
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 784

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tattttatat ttctctttt ttctgctcan agtaagatat aactttcaat tcattctatc 120
ttggctagca aaggaagaaa tcaattcttg cacagtcgat ttgttttat agatttacct 180
acatggagaa acaaaagtca atttgaaatt tcattggtac aaatatttga taattggatg 240
cttaaattatt tctagcatta aatattcata aattgcatat ggcttctttt gactcctcaa 300
aaacataggt aacaaaatac attgcanatc caatgtagat tactgggtca ttattatact 360
aatattgtta ttccaacgta gattgctana tatagtagt 399

<210> 785
<211> 333
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 785

agctnggaag gatgcttcga tggaggataa gaaatagga gagggagaga gaggggggag 60
cacgaaattg aatgaataaa agaggagag aagtggaact ttgaagtatg tatcacaaga 120
ctctcattca tcagagttac aacaagtgtt acacatgctt ctatgtatag acttggttagc 180
ttccttgaga agctttcttg agaaaacttt cttgagaagc ttgtttgata agacttcctt 240
gagaagctag agcttagctg cacacactcc tctaataaga aagctcacct ctttgagaag 300
cttccttgag atgatcgtag ataagctaga gct 333

<210> 786

<211> 200

<212> DNA

<213> Glycine max

<400> 786

gcttcacaat gacacaaggg agtgagactg tgactgagga gaggaggggt caggtctata 60
gggtgagaga ctgaggagag ggaatgaaaa tctgtatgac gtgagagtga taacaacaca 120
tgatggccct ttacatgatg gttttaataa aaccgatgtt gagtatctca ttgtcacaat 180
ggttatgaca aaaaacatct 200

<210> 787

<211> 365

<212> DNA

<213> Glycine max

<400> 787

tagtgacacg agtagccatc ccacccatgc gagtggtccat cttatcaaga tgctcaccca 60
caaagctcg aagatcgctg agctttgtga ggactctatt ccacagagaa caacaagcat 120
gccttcgtgg tggaggagat ggggtacgta catcaagaac aggtgggtggc aagtcttgtt 180
agcgaactca ctgaccatta acatcctttc gataacaaa ggaggtaaca acaccggcac 240
caatggagaa agacctttta accttgacat atgggttcac ctccaaagga acattggatt 300
gatgacgaca tagagtatca aggcggcgat acatgagagg tgcattgacc cgttatgccc 360
tatgc 365

<210> 788
 <211> 117
 <212> DNA
 <213> Glycine max

<400> 788

agcttgctcg ccacgattga cgaagggcac atgatgacga cgtagtctc tgcgtgttat 60
 caagcttttc gtcttacaga tagcctatag tttatacgga ctaccactcg ggtatatt 117

<210> 789
 <211> 115
 <212> DNA
 <213> Glycine max

<400> 789

ctataaatct aagcttaaca tcagaccctt ccaggtgtct gaactacttc acatttattt 60
 gatggtgcct atgcaggttg aaagccttgg atgatagagg tatgcctatg ttgtt 115

<210> 790
 <211> 61
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 790

agctntggta cctattgatg gataacctaat ttgttgatct aattnttagc attaacaaac 60
 t 61

<210> 791
 <211> 405
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 791

agctncacgg cagcactgtt aggattaatg caatgattgc tgaactaaa gctctacagt 60
 cgaatgagac ctggaggctc actcttcttc ctccacagaa aaccgccatt ggctgcaggt 120
 ggatttacia gatcaagtat cgcgctgatg gctcgattga aagatataaa gcacgtntag 180
 tggcataggg ctacacgcag atggaggggtc ttgattatct tgatacgttc tctcctgtag 240

catagttgac taccggtcgt cttcttcttg cccttgctgc cgtgaatcaa tggcatctgc 300
 ggcaactgga cgtaataat gctttcctcc acagacaact tgatgaagaa gtttatatgc 360
 aggttcacc gggattgacc gtttcacatc ctcaactggg atgtc 405

<210> 792
 <211> 411
 <212> DNA
 <213> Glycine max

<400> 792

tcaagaaatt gtgaaaagaa taaggtaata tacatatata tttagtacaa atagatatag 60
 ctttacacac tactgtatga ttgtgataat taaaaaagaa aaagaaaaac agttttcttc 120
 tagtaggaaa ctaaattcat acctaggtaa agttttaaga atatctaatt acctatgcaa 180
 aatttcagca accgaggcaa gatataaaca ttgtatttta aggataatta gccactgaat 240
 gtgtccttat tcttcgcaca gcaatgagga actaatccaa atgatatcag aaaatgggat 300
 aacaccagtg acaaatagac aatagtacta acagcaatca actacggaaa gttaaggaat 360
 attgtatttt aaggataatt agacacatgt gtccttattc ttcacagaac a 411

<210> 793
 <211> 502
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 793

tgatctccgc cggactaagc acgcgctgca cttagaaatt aaccgacggc tatattacct 60
 taactattga aataaaaata agtgctatat taatgacctt tggggtaaata acaaaaataac 120
 attactcaca cattttaata ttttgtgggtg gagaataaag gtggtcgcga ctatagcaac 180
 ttctagggcc ggcgctacta ctggtatagt ataaaacttc cgacccatt ggccaaaggc 240
 tcttcgctat gcgaaggat gggggaggga tggtatcgc acccttacc cttgcatatg 300
 caaagaggct ggtttcggat tccaacccat gaccaacaag tcaccaaggc acaactttac 360
 ggggtgcacca gggctcgccc tcctactact tgcatatgat acttaacaga aattgcgcca 420
 tcagctgcca gcagaattca catagagaag tattattaaa ttagatggca tcaatatata 480

cagtctgatg agtcagcttg gn

502

<210> 794
<211> 454
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 794

ttaaactcttc aaaggtagtc cgtctgaatt caattcaaatt cccttgcccta ggattcacaa 60
gcaggctgct aactcttgct gatttggcct tagcaaagca aatcttgcatt atgccgaaag 120
actntatcct tcctctataa ccactgggtga ttccaaacga gagttgaggg cagctgagtc 180
aaatggaatc cattgcccc taactttgac ctcttttggt gacttgtctt catgatcata 240
aacatttgca taaaactcct ttacaagtgc cacatctatt ctccctctt ccaaattcgc 300
gattntctta tgctagttcc tcctgtttta ctcccttga aattcctcan agtcattgaa 360
cgacaattgc acattccttt caggaatgat ctttctgccg aagatattgt ttgtgtaatg 420
atcccatgct tcangggaag aaaatttgtg tcta 454

<210> 795
<211> 306
<212> DNA
<213> Glycine max

<400> 795

agcttgtcgc tagagctgac ccatcaactg ccctaactct tttagactgg tgatccctag 60
gctcttgacc ttgacttgat agaacctctt tctaagcgaa ggcatttgac ttgatcccat 120
gttttactaa agtgaacaaa aatcgggtgcg aatcaaaaact ccaacatcta tcatgggtgg 180
aatggatgaa tgcattgaaga aatgcatatg acacatatgc aatttatgaa tacgggagcc 240
cgggaaatgg tctccttctt agatacaacg tcttggggta acaaagcgcc caacgtatgt 300
atttaa 306

<210> 796
<211> 399
<212> DNA
<213> Glycine max

<400> 796

[illegible]

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<223>      unsure at all n locations
<400>      797
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<210>	798
<211>	172
<212>	DNA
<213>	Glycine max

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gattgtcata gagtatgcta attagttata gtattatata gtcagggatg cacaaatctc 120
tccccaagca attttttcatt tgtatctgct ctcacagcac cagtagtggt gc 172

<210> 799
 <211> 106
 <212> DNA
 <213> Glycine max

<400> 799

ttggcagaga gccagaaaca ataaatgatg acgtttaagc taatattaga aagaaaaatt 60
 gcaggaagcg aagtgatcct ttttatggct acataccaaa aacccc 106

<210> 800
 <211> 372
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 800

gcacctgccg ctgcagcttg gacagtaaatt ggaatggtag ccctccatac atggtaatat 60
 tattcaacgg acccatacta tgattaaaag aactgaaac tggntctcga aatgtgatcc 120
 catttgatcg ttgcatcgag attctgaatg ttctttccaa ttctatttct tttctcactt 180
 catcttccag gtctcgttgt tgtgcaaattg tcatctcgcg agcaatgatt tctctcctga 240
 ttcttctctt ctctagctct cgccatattt tctctttctc caactctcgt tggaacactt 300
 cattgacatt gatcggcatg gccattagaa aaactctagg cactgaaaca tttctgggaa 360
 agccacctga ta 372

<210> 801
 <211> 438
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 801

nngggaataa tatntaacia ctgtagtatt gaacttaacc cacaacaagt ttctagttat 60
 atgaactaat atatgtgtca cttctaattg ttctatggct gttactttta ggtaaagact 120
 gggaagacca gcttggagca aatatcaaga gtttattgga atcaggtaag ctaaaagcta 180
 atagtcttgt ccattctttt ttcttatcca tgcaccttta tgtacttgag aatccctaaa 240
 catacatgta acaataattt ttccccatat gtaaaataac ttgacaccct cgaacttctc 300
 aaagtcattc caatttctat tcgattcgcc attggtactg gtattttcta cagattatgc 360

tcctgtcagt gttttgtaca atcaattgtc tggagttcct ctattattca tattcccaat 420
atgaatattt aaaatgaa 438

<210> 802
<211> 451
<212> DNA
<213> Glycine max

<400> 802

atcctcttag tcacctgccg catgcaagct tgaaattgac aacggaagct ctccagaatc 60
tcatatgggtg ataacttata acacgaaagt ctgattcagg cgcatagtat atctagaccc 120
tcgaaattaa acaacgaaag ctatcgagaa actcatatgg tcataaattg tcacacggaa 180
gtccgattca tgcgcataat atatcgagaa ggttggattt gaaccaccaa tgctctcgag 240
aaattcagat ggtcataact tttcaaacag aagtcgata tatgcgcata atatatcgag 300
aacgttgaaa ttgaaccacg aatgctctcg agaaattcaa attgtcataa ctgcgtcacac 360
gaaagtccga ttcaagcgca tactatatct acacgctctg aacttgacaa cgaaagctct 420
ccagaaattc atatggccat aacttgtcac a 451

<210> 803
<211> 377
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 803

ggccgccacg gagttntccg actatgctct tgtgtggtgg aacaagctac aaaaggagag 60
agcaagaaat gaagagccaa tggttgatac atggacagag atgaaaaaga tcatgaggaa 120
gcggtatgtg cgggctagtt actcaaggga cttgaaattc aagctccaaa aactaaccba 180
aggcaacaag ggggttgagg agtatttcaa ggaaatggat gtgctcatga ttcaagcaaa 240
tattgaagaa gatgaggagg taactatggc tcgatttctt aatggtttga ctaatgatat 300
ccgtgatatt gttgagctgc aggagtttgt tgaaatggat gatttgcttc ccaaagcaat 360
ccaagtggag caacaat 377

<210> 804

<211> 153
 <212> DNA
 <213> Glycine max

<400> 804

agcttggttct tgattattcc tgagttctgt aacttgctta gaacaataaa cttggccttc 60
 tcttatttgt ctttgggctt ggcgaccacg atcaacaaag tactttcggc acctactata 120
 tgttgactcg accaacggcg ttattggaaa gtt 153

<210> 805
 <211> 246
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 805

gcttcctttt aagtgcgtca ncgtttaana accgagtctc tgatgggtggg ggcgaagcct 60
 ttgatggtac cctcggcggg aagtgaatgg nggaacatcg acacttcctc attcagaata 120
 cnggccccca cactttgcaa tgggtgtgtat ttcaaggtaa tgggatataa tatccctgcc 180
 tatatgcttg cctcttggga agaaccttgg attcatgccc ctgggagtgg tcccttcaac 240
 gattca 246

<210> 806
 <211> 368
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 806

actaatgtag ttntaaaca aaaatcaatt gaggaagctt cgccaagtat cccattgaa 60
 aaacctttat tcaaaccttt caaagttagt gaaaaggcta aacgaaaaat tagggaactt 120
 agaaaaacta aatccttaat tgaaggcgta ggtgacaatc atagtgaatt actaaacaag 180
 aatggtagtt tacttaaggt cattccagat actccccaaag cctcggaaaa tacttccaaa 240
 atggtaacaa gaagtacctc caaattaatt aatattatta atgaagatag tgacccaaaac 300
 tcagataaca caactgagat aggatcagtg tcagaaaaga atataaatcc aattaatttc 360
 aaacactg 368

<210> 807
 <211> 223
 <212> DNA
 <213> Glycine max

<400> 807

agcttgagat gaggaagtgt tgaaggggtga aactttctgc ttttattggt gaccacagag 60
 tggtagctgg agatatgtcg tgggggtcac gagaccttgc ggacgtcagg tgggtgtgcta 120
 ttgccccaaa ccaagcttga ccaatcccgga cccaaccggg gcatagtcgg taagtgagaa 180
 cctgtgatgt acctaaacag gcgagctcct ggcagtcaac aga 223

<210> 808
 <211> 427
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 808

gaggtccagg aaggtttggc ggccgaagga actagttccg ctccggagta cgacagtcac 60
 cgcttttagga gcgctgtaca ccatcagcgc ttcgaggcca tcaagggatg gtcgtttctc 120
 cgggagcgac gcgtccagct cagggacgac gagtatactg atttccagga ggaaataggg 180
 cgccgacggt gggcatcact gggttactccc atggccaagt ttgatccaga aatagtcctt 240
 gagttttatg ccaatgctcg gccaacagag gagggcggtc gtgacatgag atcctgngta 300
 aggtgtcagt ggatcccgtt tgatgccgac gctatcggcc aactcctagg atatccngtg 360
 gtgttgaag agggccagga atgtatggcg cctactangc accctttgga cccagataag 420
 tncaaca 427

<210> 809
 <211> 361
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 809

agctttgata tggaaattaa gttganagta aatgatggta tgtataagac atcaatcagt 60
 gtaatgaatg cagaaagttg tactgcgctt gagtgggtag catagacaag atggccattt 120

ggtaatctaa ccgtgatggg attaatTTga tgatatgagt gaaagtttTgt taaggaggag 180
 gaaacgtgat cagtggctcc tgaatctaT atccaggagg tagagtttgc tttttcgtaa 240
 gataggatta tacctgttgc atcgttattg gaacaagata aaatggaagc gacctgtggT 300
 ttggtggatg ctgagtttcc agcagatggc tgttgTatta atgctagcca tgccttTtac 360
 t 361

<210> 810
 <211> 360
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 810

ntgcggatat ggtcttttca atgaaaggat caaagtgggt ctggaaaaag gcaaatatga 60
 tcatccttct tggatgaatg agaaaactgg ggcaaatgaa gaggatgaga atgatgaagg 120
 aacccatggt gaggctggca ttcttacaag gacaaacttt cctccagtt caaaggccac 180
 ctatttaagc ctgaaatcag aaatagaagt ggacgttggg cttttccttg agcttttgca 240
 tttttagata tttctataga gagaaaggTc caagttccaa agagttttga gagcttttTgt 300
 tgtgcgaaga ctgacagaga actgagagtg aataggaact cattctgaga catgagatga 360

<210> 811
 <211> 462
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 811

tgatgatgac aacttctgan atcaagaaac acacacacac acacacacac acacacacac 60
 acacacacac acacacactt tttcctagTc gatcactcac ataaanttcc attctcccc 120
 ttttgTTTTg aatttatgct tatcttanaa ttaagttaat tactcatgtg agttcttgat 180
 ttaatcccta tttctctcac ctttggcat caacaaaaag ccaaagtgca tatcanattt 240
 gaagtattca aatataacta aatatccata caacattcat ggaaaaaaaa tatcaaccaa 300
 atcatgaagc aagaaccatg aagcaacaat tatgaataga ttataaaatc cacatagtca 360
 aacaacatac ttaatatTng ttcanatacc ataataatat agccaaaata caaggctgaa 420

gatcagagta ctaataatat taanatagac atctaagatg ag

462

<210> 812
<211> 297
<212> DNA
<213> Glycine max

<400> 812

tggcactgca gatctgatct gcgtgaccc catccacgta cattgtgtat gctccggttg 60
agtcggtgta ccccttcttc gtgtacacaa cctcgttgct aaccctgctc ttgcattgca 120
acataatctc agcacctgca aacaatcatt aatcaaacaa tacgtgttac aaactagcaa 180
caactactaa ggtaatgttt ccattcacat taaatcattc aaaattacct tctaatatgt 240
aaattttaat atatgttcac atgttttagtc tcatgccaaa aaaatgatgt tagatct 297

<210> 813
<211> 420
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 813

gcttcatata tatataacat aacattatgt ggtaagacct aacacgccaa caaattattc 60
tcaattagtt gatgaacttc caagatccca ttctactaaa ttctcttagn tattagatac 120
ctctaaatat atgacaaagc atggatttga atatatagat aagctcatcc ttatctttgg 180
gatccacctt agaatcaata gaatattcaa aatagtgtta aggagcaacg aaaagaaact 240
aagaacaata ttaaaaggaa taaagtgtca accatctcac tcaatgaatg aatgctntga 300
caacctcaat ggatgcagac aaaataatca gaaattctag tactaatgta tcttttagcag 360
cttatgggta cttgaatgga gaataagtgt nttttctgag cttatggctc aacttgatat 420

<210> 814
<211> 432
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 814

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gacagttttt agttacattt ctttctttnt tctgctttgt attattaaat agaatttatt 120
 gttgagagga agcagagagg ggtgttatcg ttgagaggat gcagagaggg gtgtggcaaa 180
 atgacgtaag cgggttcctt ttacaatcta tttatccac aggggtctatt tttaaacaat 240
 tgtcgaagag ggtctgtttt ttaaagggtg tccaccactt ggactggcgg aacccttgct 300
 tgcataaatt cccatgcatg catggttctg ccagtgggtg tagcgtaaata caacccaaag 360
 ggtttcgcca atcccactag tgagataccc catgccattc atcccacat ccatcaacgc 420
 catccatcca tg 432

<210> 815
 <211> 429
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 815

ctaaaccttg aaaacttggtg ctattcattc tttcatctc ttctcccttt gccaaanaga 60
 attcgccaag gacctaaccg cctgaattct tttgtgtctc ttttctccct tttccaaaag 120
 aacaaaggac taaccgcctg aattcttttag tgtctccctt ctcccttgct aaagaattca 180
 aaacgacaca gtctgaaaat tcttttgatt cttcccatc cctaatacaa aagtgttcaa 240
 aggactaacc gcctgagaat tcttttgat ccccatcac aatgtatcan aggtttaaca 300
 gcctgagatc tttgtctaaa cacattggag ggtacatcct ttgtggtaca agtagatggt 360
 acatctactt gtgtttgact gagaacaaga gaangtacat ctcttggtga tctgttctag 420
 tggagggtgta 429

<210> 816
 <211> 529
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 816

cgcccatcat gattgaaacg ccatctatnt angtgtgact acagataact caagctgtac 60
 ggtattgcgg gcaagttatg tgctgaagat ntattntata aaattaatta attaattgat 120
 gtgagttaaa tttgaaataa aataaagaaa caggacatcc atttaccaga acgtgtccca 180

attaatTTTT gttactatTT ttaattacat aaatgatata tatatatata tatatatata 240
tatatatata tatatatata tatatatata taattggcat gaatcgaaaa tattctTTTT 300
ttctTTTTta tcgtaatTTT aaactgtaaa aaagtaggac atTTTTttcc tatagcgtga 360
tacatacaaa gttaacagtt aacaattgtg tntatataat atataaatat atattaatta 420
ataatatata ttacttctag taaaaataaa ataataact gcttaataga tntgatgatt 480
aattccatcg attatatact aatctntnta atataaaaaa ataaaactn 529

<210> 817
<211> 501
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 817

agtcacctgc ggcattgcaag cttctgggtg ctgcgatgat cgaccagaac tgttccctga 60
actcatccaa gtcctcaaag gcttgagact cacagcagtt aaagctgaca tagccagtgt 120
tggtggcaga atcaaaagca tattggtgct ttgttctaag gatagagaag acagtgtttg 180
ccttgccact ctcaaacagt ccttcaaate tgctgtcacc aaaattgctt catcatccat 240
ggcttctagn tgtcccgcta gaagtaagag gcagagattc ttcttgcctt ctactgcct 300
acagttaatt atttattgca aanaatattt ttttccccac tattcattgc agtatggggc 360
aattatttgc tctattntca atatatatat atatatatat agactcccca ttaggaagat 420
aaaatcatga aatattagtt ctgtgcaacc aattaaggca tagttaaatt ganaggaaaag 480
gacgcacagc atagtatgag t 501

<210> 818
<211> 447
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 818

cgcttaatta tcttttagagg aggataccgt ccaatggagg ataaaaaat gatgttgata 60
atcttttccct tttttacatt tgaaggtatt aaaaaaagt gtcttcatga cattattaag 120
aaattaaaaa aattattaaa aatcatataa tgcatttaaa aatcattgga gaagacattt 180

ttatgcattt caatgaaaat atttttcttt ttattttttt taaccgctgt tcttaaaaca 240
 ttagttaata ttaccttaa aaaatgctaa caaaagatat taatgccttg ttaattagaa 300
 gtattaaaat aaaagattga ttctttntaa aatgtatata ttgtttataa tattttattg 360
 ataatacttg taattagtgt cacaattaaa agtagctttt tataaaaatt gaaaatgaaa 420
 acttccattt gaagtcagca atgttat 447

<210> 819
 <211> 429
 <212> DNA
 <213> Glycine max

<400> 819

ttcgttggcg aaaggatcta tgtgggtctg agaagaggca aatttgatta tcttgcctta 60
 atgaatagga agcctgcggc aaatggagag aatgagaagg agggaggaac ccatgttgtg 120
 actgctgtcc caacacgacc aaatttccta ctagctcaac aatatcaata cttatccaat 180
 atcagccctt ctcattaccc accaccctat caaccaagaa cactcaatca ttcacaaagg 240
 tcatacctaa atcagccaca aagcccgct accgcatatc caataccaaa caccaccctt 300
 aacacaaacc acaataccaa ccggtgaatg gaatttctag aaaagaagcc tgtagaattc 360
 accccaattc tgggtgtgta tgctaactta actccatatc tactcaataa tgcaatggta 420
 gctataatc 429

<210> 820
 <211> 466
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 820

nttccctcat tctcacgttg cttttcttct ccccttctca tccaccattg ttgccatta 60
 aagctccaaa ctttgctcac catttctact ccaaatacaca aaaggaagcc attttcggag 120
 tcgtgaagcg cacctctacg ttgtgggact tcaaatttca ggtttgggta gacttcttct 180
 cacataaatt ttgtgggtat tgggtctttg ggagatatga tgggtagttc tactaggttt 240
 atgccttatg gtagttatgt gtgaaggaat ttgttgaaag catgctaaac tcgtcatgtt 300
 tgatgtgagc caaatatacc cattctgttt tagggtttta taatgatgct ttgtgatgct 360

tgtgtgctga aatcattggt agaaaactgg tagagatgat ggggagagtt aacctanggt 420
 taaaagtgag aatggtagtg atgtgagtgg aaaagtgagg ttttga 466

<210> 821
 <211> 483
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 821

gcttcacact tgataatgga gacacatgaa cagtgtctatg taatgacatt catggtgctc 60
 caaaciaaagg tggagtatgg aggattgcct tgagggtccg cacttaggca atcatgaaac 120
 tcaactccaa actcgaaagt ggaggacaca tgaacagccc taagcaataa cattcatgtg 180
 gctccggaat aggatgagaa tggaggattg ccttgagggt cctctcttaa gcaatcatgg 240
 aacacaactc caaactcgaa agtggaggac acatgaacag ccctaagcaa taacattcat 300
 gtggctccga agcangatga gaatggagga ttgcctcgag ggtcctctct tatgaaatca 360
 tgaaactcaa ctccacactc gaaagtggag aacacatgaa cagccctaag caataacatt 420
 catgtggctc tggaacagga tgagaatgga ggaatgcctn gagggtcctt ctttaagctat 480
 cat 483

<210> 822
 <211> 412
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 822

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 tgccaaacaa agtcaggtaa acgataactc gcctgtgctt tttcttccat gctatatgta 120
 gcaaagtcac tgatccagtc atgtttgatg atttggaaaa tgaggccgca attatactgt 180
 gccagttgga gatgtatttt cccctgtctt tctttgacat catgactcac ttgattgtgc 240
 atctgggtcag agaaatcaaa tgttgtggtc ctgtttatct acggtggatg taccgggttg 300
 agcgatacat gaagatctta aaaggggtata caaagaatct atatcggtca gaaacatcta 360
 nttgtgagag gtacattgca gaagaagcca ttgaattttg ttcagaatac tt 412

<210> 823
 <211> 453
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 823

agctntagga gaaaccatat aaactaaggt agttcctana caaaaaacaa ttgaggaaac 60
 ttgcgaaga atccccattg aaaaaccttt attcaaact ttcaaagtta gtgagaaggc 120
 taaaagaaaa attaggaac ttagaaaaac taaatcctta attgaaggcg taggtgacaa 180
 ccatagtga ttactaaaca agattggtag ttacttaaa gtcattccag atacccccca 240
 agcctcggaa aataacttcca aaatggtaac aagaagtacc tccaaattaa tcaatgttat 300
 taatgaagat agtggccaaa actcagataa cacaactgag ataggatcag tgtcagagaa 360
 gaatataaat ccaattaatt ccaaacactg gagaacaacc tccatattat attatcaacg 420
 tccaactggc cctgaccttc tattagagga aag 453

<210> 824
 <211> 457
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 824

ngaaacaact cgttacataa tttgatatgg cttgggcaac agatttcata agtatttctg 60
 taccattct agaaagagat tgctccatcc acttggtcaa tttttattac acactcttct 120
 aaatgctaga gaaaatcatc ttaaactttc ccaataaagt ggtagcccaa gatatttctg 180
 atgaaaattg taacgacctg tcttgctggt atgatcac cactctaaag tacgtaaatt 240
 ntaattttta aatgaaaatt tcattaattt gcttatgaaa aatgagagta aatttttctg 300
 gatatagatt caccaaaca cgcacaatta tttaaagaa atatatatat atatatatat 360
 atatatatat atatatatat atatatatat atatatatat atatatnaac ttagccacac 420
 tcacataata gaaaagtaaa ttagttcata catatag 457

<210> 825
 <211> 479

<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 825

agctngaaca cctattagta tttatatttc ttaaataata aatgtatgac atcaactcgt 60
cattggctcc caggttagtg gattaagcaa aatagaccaa tacaaactca cgggttaagt 120
cacctaaccc attgatccaa agtttacatt gtcacctcta cattagtgcac tntttgttgc 180
ctttgtttcc ttttaagcttt ntgtgtataa aaatatattt tttcttgtgt gaaatatttg 240
tttggaattc agttttaact atataataaa attgatgggt aagtttaata tatatttaaa 300
cagtcttgat catttgatta tgaggacttg gataaaatat atattcttca aagttttgtt 360
aatataactt ggtaaataata attctaattt tataaactat ganaaaatac aaaaggtaga 420
tgaattcaag ctcaacacaa tagaacaagt accaacanat actatcatac atttgacat 479

<210> 826
<211> 430
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 826

tctgtctgca gtagcaccac caccagccat gttaattaat ttgctctcaa acaaccaatt 60
aattctctag cctcaacgct ggtacgagct taattagtat attaagaatg ctattattag 120
taatataat tcagttctat agagaatgat gttttgtcat atgcttacag accgtaatgg 180
tattgctctt gcgggaacca catacactct agctagaaaa caagacatac atagttaatt 240
aattaataat gttaaagccg gcccttgga gacaatttac atgcttaagt ttcacggggt 300
taggtctaaa taatgccatt aaattatttn ttttgtttgg aatataattt atttattatc 360
atttaagttg caacaagcct tggctggttac ttgtttgttc tattatgcgt ggagtcttat 420
tcatcgaacg 430

<210> 827
<211> 309
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 827

agcttntgac cggtgacaa gcaacaatct aagttataat atctacacca caagctgtat 60
cattgtaata acttactcat attcttcaca tttggacaga cgataatatac tacaccacca 120
gctgtcctcg aataacctga attgaaacag aataaacagc agtaatggaa taaaccccat 180
cgaagcatct gcgtaagaaa ccacttgga ggtgtgatca aatatcaaaa aggaggctgc 240
tgtaaaagcc acaaccatat catcaacata gataaaaagt aacacggtcc caaaagtcaa 300
gctcggatg 309

<210> 828

<211> 222

<212> DNA

<213> Glycine max

<400> 828

cttctgaacc ttcctaagga aatccttgat aggattgaga ctcaggaaga tatgaactat 60
attgaagggtt ggtagcaaat ctttacttct acattaacgc ttctcattat gtataggata 120
cgagtgaaaa ttcatatctg ggatggggta tttcggatga atctttcttg ccggcggttcg 180
tcattcgatg ccccatatat accctcatta aagggccaca ac 222

<210> 829

<211> 406

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 829

cccgtgagt tctttgttat caccganact ctaggaggag gccaaagaaac tactgtgtaa 60
gattttcttt atttccttgn tagtggttctt gatttgtgaa tctcacttaa attttgagct 120
taatatgtgg catgcattgt gaatcacatt tttaatcttt atcagctaag ttgagttggt 180
tatgtatggt gtagggcctt tcaaggagaa acgaagcaat gagcttatat tctaatagct 240
canaatcaca tataattctc acatttgtca ttgagtcttt gtgtaaggga ctgtcaaatt 300
ttgtaattct acctaacatt accagcagtt gtgtatggaa attgtntggt tcctaagatt 360
caagccaggt ttatatcttc tctgttagtt ctattgctaa acatga 406

<210> 830
 <211> 399
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 830

 ggggtgtacgt aaaataatta gcactaaagc ttatattatt ntatgtattt atcacaatgt 60
 gcaaatgaga tgattgtgtc taactttttc cataacatat aggctgatg ttttatttta 120
 gaagtttata atctgatctt cagaatttga aattttttct ttacattctt atcaaaggaa 180
 cgtcgtttta atcaggtaga gtacaattat aaacatataa attacaaaaa ttaaataataa 240
 cattntatgg tttatgatga tgagtgagat gtggaatgg gagagcaata aggaaagcat 300
 tntggatcca canaacagga catgaagctt cttgcaagtt ttcaaataat cccttcacga 360
 atcggtttgg gatctttgca aatttcttgt ctcanatcc 399

<210> 831
 <211> 440
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 831

 tgcccaagtc agcaacatac aactgtgta tattgtggat tctacaataa aaggtaactt 60
 tgaatacttc acagtcatga agtgtctacc aatctctgtt aagatagttg ttacaacttt 120
 atttaagcaa agaaatagct cctttcctta aacacaaaaa cagaagcaac aaaagttagc 180
 aaagcaagaa atacaaccgc gaaccataaa atagtgtctt tatagattca tgattcttaa 240
 tattttcagg tgtgttaaaa acgatttttc tgaaaaattt gtcacaaatc aagcttgaac 300
 cagagaatca ccacgcgcag ataattaact gcacattagg tgccgtagct atcgagataa 360
 agaaaagtag aggctattgc tcgagaagaa nagaggaaaa gcttatgaac aagaaagcan 420
 ataaattgag caaatgatc 440

<210> 832
 <211> 313
 <212> DNA
 <213> Glycine max

 <400> 832

<211> 254
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 835

tgagatgagg aagtgtagaa gggtgaaact tcttgctttt attcggttgac cacagagtgg 60
 tacctggaga tatgtcgagg nggtcaggag accttgnnga cgtcagggtgg ggtgctattg 120
 cccaaaacca agcttgacca atcccgaccc aaccgggca tagtcgggtca gtgagaacct 180
 gtgatgtacc taaacaggcg agctcctggc agtcaacaga taaaaggaac aaagaccaca 240
 aagcatggag gctt 254

<210> 836
 <211> 402
 <212> DNA
 <213> Glycine max

<400> 836

tgcagcttgg ttaatatag gccttgataa tcaataaagc ctattggaaa ttaatatgta 60
 aaaggtagtg actaaaaatg caaaattaca ctttagtttt tcataagcat aacatccaat 120
 acaactcata agtttataat tagtcacata agtttttcat aacatatcac aagtcacaac 180
 taaaataaaa gaaaacagtc caagtgtgca attatagaat ctagaattct tgatatttag 240
 actagcacca aatcgcta at tttttccaaa taaaaacaca aaggtagtaa tagagatggg 300
 gaggggcact tgatgtgaga gaaaagctta tctctgctaa gcctcaggcc aaagtccaag 360
 ggtgaatatg gagagatcaa aagttacaga gcaatgggta ta 402

<210> 837
 <211> 433
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 837

tgatttagtt ntcgctgacg aaaggatcga agtgggtctg ataagaggaa aatttaatta 60
 tcttgcttgc tgcttgacg aatgagaaaa ctggggcaaa tgaagagggt gagaatgagg 120
 aaggaacca tggtgtggct gccattccta catggacaaa cttcccttca gcccaataat 180

gtcatcgctc agccaatatc gacccttctc attaccacc acccagtcac ccacaaaggt 240
catccctaaa tcaaccacaa aaccaccta ccacacaacc aatgctaaac accaccttta 300
gcacaaacca aaacaccaac caaggaaggg aatttgcagc aaaaagcctg tagaactcac 360
cccaattctg gtgtcctatg ctaacttgct cctttatcta cttgataatg caatggtagc 420
gatcacccct act 433

<210> 838
<211> 111
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 838

agaggtaa at gacatatgat catttctcct acttgcgtt gcacttttgg ctatttttga 60
tacctaaagn tttcattncg aatttaatta cacgatgatg atgctggatg a 111

<210> 839
<211> 402
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 839

agctngcgaa taattaacac agaatngtac aaaattctta tgatacatga tcaaatttat 60
caaaaaanat aataatgacc cctgaagcta tcttggtgac agtgacaata agttgagcct 120
tgtgcagcaa aacttagtgt tgagtgaagg atgacttggt gcttgatgaat tgacttaacc 180
agtttttgac agctttacct ttggcaatga agcagccatt gttcttctta tcaaccttca 240
acggactccc cattttgacc atcatcttct gaaccaacac tctcacgggc gactctctc 300
tgacactgct atgctttccc aaccacana ccaccgtaaa ttccgcagga actccataat 360
tggaatcatt caaccttctt ccatctctc aaccaaagca ca 402

<210> 840
<211> 116
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 840

ttctttttgt gggaatatga gtccgatcac acaagatggc atgatcactt gcagccatgt 120
 tctcaataag ctccatggct tcttctgggg tcttcaattt aatttttccc ccagcagaag 180
 catcaaataa ctgcttggac tgtggcctta acccatctat aaaaatggtg aactgaattg 240
 gttctgaaaa tccgtgagtc agtgttttcc gcagcaagct atggaatcgt tcaagtgcct 300
 cgcttagaga ctcatccaga aactggtgaa atgaaaagat ggntgctttt ccttctactg 360
 tcttagactc ggngaaatat ttcttcaaac atttctccac caattcatcc catgtcttga 420
 gactgtntcc cttaaa 436

<210> 844
 <211> 434
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 844

agcttcatga ctagacatga cttctatgac aaaactacaa taggtggaca agtcgctcta 60
 gatttgtgag gttttcttct actttaatat ttttgtaaga attttatgat ttaggtttca 120
 gccacaaaaa ataacaagac aaaactcana tcatttggtc atgagtgtat gaaattcttt 180
 tagcctatta ttgatttga gtcaaatctt tcatgttaat tagtccttaa catgttcatg 240
 caaatgctt agagagtctt tgattgtgaa cctttgcttg aacttttatg cttccttatg 300
 attgctgcta ttgtgaatat gagtcttggt gattgaattg ctggctgaaa tgttgatcct 360
 aagtgaatat tgaactccta taactgtcgt aaacagtcct agtgagttca acatacatat 420
 gaagggtgaa agta 434

<210> 845
 <211> 376
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 845

ngaagctcaa ggaaaagctt gaagaagtn tggctattac atgcccaact ctcttaagt 60
 gcatttgtat tgggtgttat ctnggtgtt tcttcttagt acatttgata ttgtattgc 120
 atcatgcac atcatggttt gtgtgaagaa aagtttctaa gttagaaaaa tttcttcaga 180

ggcaaaaaca ctattttaat cgattacaac cttattgtaa tcaattacga caagctgtct 240
gaagcttata gagttgagtc tcgtatcaaa ttaatcgatt acagctatct cacaattgat 300
tacattattg ttogagacaa tgactgattt attcaagagt ctctgcttta atcgattact 360
tctttctcgt ttaagt 376

<210> 846
<211> 415
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 846

agagatacaa tctactcttg gatttgcatt aaaanaaaat ctttgcattt atctttcttc 60
tgctccctgc ctgaaaacat gcatgtatag cctggtaatt ggtcaatgtc tttgctaaag 120
attatctggg ctggtgactc gaagcttttc taatttattc aataagatgc atgaataatt 180
taatcataaa tcataaatc catacatgat gtaattattc atgtatctta ttgaatatat 240
agatcttatg gttatttcat ataattagtt aattaactgg tgattatttt ctgaccaagc 300
ctgggtgatta tttcatacgc ttacgtaatt aactgattct gttttatatt ttatttatta 360
attcttcata atggagatga antctacaca attcatgttt gaacacagga tgcatt 415

<210> 847
<211> 455
<212> DNA
<213> Glycine max
<400> 847

tctttgagaa aacttccttg agaagctaga gcttagctac actcaccct ctaataacta 60
agctcacctc cttgagaagc cttcttgaga agattcctat agaagctaga gcttaggtac 120
acacacctct ctaatagcta agctcacctc cttgagatga aaagctagag cttagctaca 180
caaccctat aatagctaag ctcaccccca tgacaaaata catgaaaata caaaaaattc 240
cctactacaa agactactca aaatgtctcg aaatacaagg ctaaaaccct atactactag 300
aatggccaaa atacaaggcc caaacgaagg aaaaacctat tctaattttt acaaagataa 360
gcgggctcat atttagccca tgggctcaaa atctacccta aggctcatga gaaccctagg 420

gccttcctt ggatctctgg cccaatctac ttgga

455

<210> 848
<211> 349
<212> DNA
<213> Glycine max

<400> 848

gcgagctctg accactgttc ttctttccg cgatgcttct tttcatgtgc gccggagtgg 60
gcttatagcc taaaccatac ttcccacgat tcccttgggt ttttatcaga ctagttatgc 120
cgccattgtc tttgcctaaa cccatcccgg gttcataacc ggtccccaac ataactcggg 180
ccatcattac cgccgcatct gacagacaat gttgcccaaa gagggaatcc acggaggaaa 240
tgctgaccac ctcaaagac tggaaagcgg tttctaacga ttcttctgcg gcttccacat 300
aaggcatgga ggatgggcag cttaccaaga tatcttctc gcctgacac 349

<210> 849
<211> 106
<212> DNA
<213> Glycine max

<400> 849

gctgatgagt actattgtgg agatgattga ccattctcaa agccaaggaa catatggatg 60
cgtatattga aatgatgaga tcattatcca acaacatgaa tttatc 106

<210> 850
<211> 358
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 850

agctttgcc aataaacaag ttctctattg ttgatactca gatcattcca ttttaattnt 60
aaatacttgg cgacccgatg cgcttgccgg tatatcactt ctgctttgat gtaagtcttt 120
gtaaatttaa gaaaaaggaa ctgtgtgggg agacgaacag taccacattg catttgagag 180
ttgaggtcag gtacatatat catactaagc atgagtgatt gaaactatgg acgaatgatg 240
actactctgt gagtgtatgt tggactaatg gatggttgcg tatgtttatg ggatctgata 300
atgttttctt actaattatt cgagttttgt attaaacttct tttataataa actcaccc 358

<210> 851
 <211> 435
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 851

tatccttatg gcttgccttc ggacttcact ccccgtagca ccccggaaga tttaagccaa 60
 gccctactt tgcaggggca gctccacact tatgacgact atcccgggca agacgatgag 120
 gaaggagata cccatctcgg tcccctgctc cacctcaaag atctgtcccc ccatgaacta 180
 ccccaaccaa acatagtccg ccatatcccg acttcaccca cactcgtaaa agaactctgtt 240
 cccttcgtgg aacataaggg aaagattgag gcgcttgaag agagggttgag agcagtcgag 300
 ggcctcgaaa attaccatt cttggatcta gcggacttat gtctcgtagc caatatcgtc 360
 attcctccca agttcaaagt accggacttt gataagtaca aagggatgac atgtccgana 420
 gggcatcttc ggatg 435

<210> 852
 <211> 187
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 852

agcttgaata ggtgtcgcgt tctactgaac tatatatatt ntgggtgntt tgcaatgttg 60
 ctatcctaga tttgntatct tctctcttgg atatttttagc gtcgacatag tgtaatatac 120
 aggtatgagc ttatcaagat gaatcatttt aggggattat cacttggcat tgttcagcta 180
 ttctcta 187

<210> 853
 <211> 333
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 853

nttaggtgga gcatattana ataaactaag ttcatagaaa ggtaataaga caccttcata 60

agcccaaatg agatcttcta aacaccctct ttatccatat aactctcaaa gtattactta 120
 acaaaatttc aatgaatcct tcaaataatgg gaactgtcat ttcaatcaaa gaaaccaaca 180
 aagtccacca taaaaaagct gtcttcactc tttagactgt ttacaaaatt cactagaaaa 240
 tattctacga agaagatatc atcatatgat gtaaagcgtg ctagttaact atgcaatgac 300
 acaagccgtc caaatgcacc cattttaaga tct 333

<210> 854
 <211> 315
 <212> DNA
 <213> Glycine max

<400> 854
 gatgatgcag atgggtttgt agctacctca tgcactcctc taatgactat ggcattcattt 60
 ctggcactaa actgctggga gttggaggcc atcttctcaa ttaaatttct ggcttcagca 120
 ggagtcattgt ctccaagggc tccaccactg gcagcatcta tcatacttct cttcatatta 180
 ctgagtcctt cataaaagta ttggagaaga cgctgttctg aaatctgatg gtggggggcaa 240
 ctggcacata gtttcttaaa tctctcccag tactcataca ggctctctcc actgagttgt 300
 ctaatacctg agata 315

<210> 855
 <211> 303
 <212> DNA
 <213> Glycine max

<400> 855
 tcagaagaaa gtgatgaggt acaagctcta aaggcagagc ttgaaagagc ccgagtagtc 60
 gaagagaagt tcaagtccat agccatcaaa gtctgaaaag agtatgatga actaaggggac 120
 gtcaatatgg ccaccgctga agccttggaa cgagaaacca agaaggcccg aaaggaagaa 180
 cacgtgcaag caaagttttg aggggcttta tatggcagca atagtttagct caagctccta 240
 agaggtgaaa ggaatcatca ctgggttaaag gcatgatctt gaaagacgag cttaaaggctt 300
 acc 303

<210> 856
 <211> 415
 <212> DNA

<213> Glycine max

<400> 856

cccatcacat gtggtactag gtggcggtcg ggcgatggg cacaacaagt tttccacatc 60
cacaatgcgc gcataaacc accatccct gttgccacc tccatctgag ctcacgtatt 120
cccacgtagc ccatatcctc gtttctctca acaccgggtc cccatcaatc ctcccaagct 180
tccacaacat ccaatcaaaa caacattcaa acagcacaag ctatcacagc caagcaaaac 240
aggacaaagg cagaaaactc tgctcaacac accaaccaaa atcacagctt ttctcactca 300
aagacccag taacaatttc ttcgatccaa ttcgttaacc gttggatcga ctccaaaatt 360
ttactggaag tctatagtgc ataagcctac attgtgaacc gtgggatcta ctaac 415

<210> 857

<211> 591

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 857

agggcacgac ggtnaattga tgcacatctat tangcgacac tatagatact caagctngta 60
atcatgtgac accctctacc cctcacatgt atactaatat atgaataaaa ttcaaattatt 120
aattaaaagt atttttaaaa cattnttttt tccgaaacaa gtctttcaaa ggggaaaaag 180
gtcacattc attttcttct acatcatatt caaacttgct caaataaata ataaagttaat 240
ctcgtctcan acaaggctgt cttaaacttca tacaattaat atagaactta taccctaattg 300
tcacatccta tcatagtgtt gtgttcctgt gtcctctagc atgaggttct tcatagtcatt 360
ccacctattc atctgtttcc ccgaacacaa gttcaagatc atcacaggat ccanacacaa 420
caacacacag ggagtgagtc atcacattca tagctaatag agagacaaga caattaaata 480
tagatattat ataaatgaga taccacttgc ttaaacaatag ctcacgtaat ntcaccactn 540
tgtcattcan naatcacttt tcaatcatca atcacattac acaagaatcc n 591

<210> 858

<211> 416

<212> DNA

<213> Glycine max

<400> 858

gtgtgattcc tttctttttc ttatcattct cctcatgttg attcagtctc attagttcca 60
 tttcgtgttc ctataacttt ccaaataaag ttgcaagaga catgttagaa agatcccttg 120
 attctgtaat agttgttacc tttggttgtc attccctact taaacatctt agaactttat 180
 taataagatc ctcatggga aatatctttc ctaatgatgc aagatgattt actatgtgtg 240
 tgaatctctt ttgcatatca tgtatagttt catttggtt cattctaaac aattcatatt 300
 catgggttaa ggtattttatt ctagaacctt ttacatctat tgttccttca tgggttactt 360
 gtaaggatc ccacatatct tttgcactct tgcaatttga tactctaaag tattca 416

<210> 859
 <211> 487
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 859

catgcgcta tntccttacg aacgttctct ngcacaagac atttagaccg aaaaatgcac 60
 ccatatacaa tcaaggcagt ttcgttacct agattattta cacgtacctc caagggtgat 120
 ttgttactta catcacacac atctccttgg ctaaattcac atacatgcat actcaaagca 180
 ttttggggca ccaaaaattg cacctgtgca catcttggca tttctaatac ctatacatat 240
 gcaaacttca tgatgaatct tgactatcta cacaataagg tgctacattt catgctcttt 300
 tttcaagtn ttgctaccta aagccgcatg ccaattcaag catattttcc tttgctgact 360
 aanatngtat tcaaattaaa aggtatatan ctttttgtaa tatagtttct tcacataaca 420
 tgcaacatat ttatatatat ttttctgtga gacatcttga ctaccaacaa tatatatata 480
 tacattc 487

<210> 860
 <211> 502
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 860

cgagccncng nnnntgtagc atctcgaccg cgatccttaa gcacctgggc tgcagcttga 60
 ccaacagaga gccataaagt ttgtctagga agtctaagcc tatagtgcct ggaggcgttg 120

cactctcata ccgtatgcca agagcttgaa gcctttgctt ggaagacgcc atagtggatt 180
 tcatcctcta ttctttaccc gatcactatg atagctctgc aggtacgcat gtattgtgca 240
 tacctatggt attgtagctc tggcagcagc ttcttcacta catgcgggac tgcccacctc 300
 agacataggg cactgagtata ccactgttga tcatatggga ctccctctct tttttaatcg 360
 cgatggcatc ccactccgct gctcgtttat gacatgataa gacagctgct tgcattcant 420
 gtttaccgta atagaagagc acaaccctat tccagcgagg cacatattat tacttgccca 480
 ccatgggtct gaattcttgt cn 502

<210> 861
 <211> 311
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 861

tgagctctnt cagctacaca tggcgggtgac ttgtagagga gtgtgatgga cgaacctgca 60
 cccgacgata aactgacaa gagcttatat tctccatctt ggacaagata tggcagactg 120
 gtggcaagtc tactatcttc catcagcctt ggaacaactg tgatcgatgat cacatatcag 180
 ctaaaacttg atgggtatgc aagccatact ctactgtgc ttgaatggta acgagcagcc 240
 caatgacact gtgacccaca tgtttctcca catgcgttac atcaatgcag tgtctaacgt 300
 caagatcaca c 311

<210> 862
 <211> 80
 <212> DNA
 <213> Glycine max
 <400> 862

tcctaacgat ttctaattat gtggggccatt aagtctatca tatgctgaca atagccgaga 60
 agcccatgaa tctcttcggg 80

<210> 863
 <211> 440
 <212> DNA
 <213> Glycine max

<400> 863

cggagaggat gcttcactgg aggagaagac agagggagag atagatagag gcgggagcat 60
gaaattgaac gatgataaac ggagagaagt tgaactatga gttgtgtctc acaagactct 120
cattcatcaa agatacaaca tgtgttacac atgtatctat attatagact atgtagcttc 180
cttgagaagc tttcttgaga caacttcctt gagaagcttc tatgagaaaa cttccttgag 240
aagctagagc ttagctacac ataccctct aataactaag cttacctact tgagaagctt 300
ccttgaatag attcctaattg aagctagagc ttatctacac acacctctct aatatctatg 360
ctcacctcct tgagatgaga agctagaact tatctacaca caccctataa tagctaagct 420
tacctccatg acacattaca 440

<210> 864

<211> 566

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 864

nttgacgcat cttgangcga cactatgata ctacagctggg tatgtcaggc caaatatggg 60
tggagtgaca aaagctntaa tttactgctt gaagtagtgc acgatctgct tccagaggan 120
aacacgttgc ctaaaagcta ctatttgccc aagaagatac tatgtccgat gggatatggag 180
tatcagaaga ttcattgcttg ccctaattgat tgcatactgt acagacatga atttgaagaa 240
atgtccaaat gccctagggtg tggngcatca cggtacaagg tgaaggatga taaggagtgc 300
agttctgatg aaaactcana gaagggtcct ccagcgaagg tgttgtggta tcttcccatc 360
attccaaggt ttaagcatct ttntgctaatt gaagacaacg canaagacct tacctggaat 420
gcanatggga gaaactctga tggaatggtc tatcatccgg ctgatntcct ctagtggaag 480
aagattgatg gtttgtattc ggatttcaga aaagaggcaa gaaatcttag gcttggacta 540
gccagtgatg gaatgaatca tatggn 566

<210> 865

<211> 441

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 865

ataacatcca agaaatttca acatccaaac atcatgaact atcaaaaacta agcaaaaaca 60
gggcagagggc agaaaactct gcccaaaaaca caaaccaata ccacaacttt tcttattcaa 120
ataccccaat cacattcttt ttgttccaat tcattcacgg ttggatcgac tcaaaaattt 180
tactggaggt ccctagtaca taattctaaa ttttgaccgt tgggatctcc tagaaaacgt 240
ccagaacca atctgtacta ctctttccac aaccagcaaa tacacatcat tttctgcatg 300
caciaagcca aaattctgct gcacatttca acagcaaaac tctgcataat agtgcaaaat 360
ttcgaaatca cacttgccct tgtcctaatt tgcccaaatn gaatcctaca agtcctaaat 420
catgtataaa tcatgtctaa a 441

<210> 866

<211> 318

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 866

ataaagaggg tgaggatgag ggagaaaccc atgctgtgac tgccattcct gtacggccaa 60
gtttcccacc aaccaacaa tatctttact cagccaataa caaaccttct tcttaccac 120
caccagtta tccacaaagg ccacccctaa atctaccaca aagtctgtct accgcacttc 180
caatgacgaa caccaccttt agcacaacc anaaacacca accaagaagt gaatattgca 240
gcgagaaagc ctgtagaatt caccccaatt ccagtgtcct atgctgactt gctcccatat 300
ctacttgata attcaatg 318

<210> 867

<211> 471

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 867

agctntgcat ggatgccaca cgtgaatctt ctgtatcatc catcatttct ttcattcaca 60
acccaacaa gatgtagtac actagagtta tgggcaaagc aatcagcatc ccaaaaataa 120
cactgcattg ggttaaagaa gaagctccat tggtaggat taaaagagaa agtaattgaa 180

tgaaaaggaa aaaggaatgg aggggagaga aagttctaata tgagccatat aagaattaga 240
 tttgaatact cacgctgtgc tgagaatatc aggatgtaca ttatattcct tagcaaagac 300
 aaatgggaca attccttggg gaagagctgc ctatattttt ttgacattca gttgccaaac 360
 aaacaggaca agcaaacaaa caaacatgtc aactgcagtt tcaaatacctt ggtgtacaat 420
 cacaacatg atcattingaa tctcactac tagtactaag atcttcaatt t 471

<210> 868
 <211> 338
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 868

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 gttacacacc cctctaataa ctaagctcac ctccttgaga agttccttga gaaacttcct 120
 tgagaagctt ccttgataaa cttccttgag aagcttcctt gagaatattc ctagagaagt 180
 tagagcttat gtacacacac cctctaata gctaacttca ccttcttgag atgagaagct 240
 agaagttagc tatacacctc ctataatagc taagttcact cccatgccaa aatacatgag 300
 aatacaaaaa cattcctact acanagaact actcaaat 338

<210> 869
 <211> 375
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 869

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 gagatgcagc ggaagacaaa ggaaaagagg tgagaggagg cgccatccac tagggaataa 120
 gccatggaag aaggagcttc accaccaaga tgagccttgg ataagaagct tggagaggat 180
 gcttcaatga aggaaaagaa agagggagag aaagagggag ggggggagcac gaaattgaag 240
 gaagaaaaag ggagagaaaag agggaggggg gagcacgaaa ttgaaggaag aaaaagggag 300
 agaagttgaa cnttgagttg tgtctcacia gactctcatt catcanagtt acaacaagt 360
 ttacacatgc ttcta 375

<210> 870
 <211> 457
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 870

ntttagaaaa aatcttataa gttagtacat acctaaatag ttctttaaga tgactatatt 60
 atgcttaaat aatctttaga aatttcaact caataacccc taaagtcag atattcaatt 120
 aaaactatca attatntaa aagagaatgt caacatttgt gatattcaat tgagactttt 180
 cacaactaat aaaaagggtat tcttctatta aaaaatatat aaattttgat taattatttt 240
 ctagagtga ttttgtgtca attctttagt atgatgtata aattctagac tcatccaaca 300
 atttcaccaa aactttcctc atttctgttg aaacatactt aatatgaagt tntgatgatg 360
 tcacaagata agcggttctc aagtttaatc caagttaaga actcagaaat tcaagataaa 420
 tgaagaagta gtccttaaga gtcttagaaa gcattct 457

<210> 871
 <211> 417
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 871

gactaaacat tcattgggta tttatttgta ttcattatgc gatataattc gctgtaaccc 60
 gtcactaacc aattaatatt atcaactact cgtttggtta agcaaggaaa ttgttggtcc 120
 aacaaaaatc atttacggt acagcatata tcattgtcat aattgacaac acataatgac 180
 atgcatgctg gttacagttt gagcgtgaca acacattggt ngacttcagt acacattttg 240
 aaactagcag tcgctcaaca acacattggt tgacttgact acacattagc gacaacacat 300
 tggctgactt gactacacat ttacgctgt ctatttggtg tgaaacanag ttaaacaag 360
 gtcggtcac aaccatctat atatatggca gactangcta ctaaatcaca cattatc 417

<210> 872
 <211> 412
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 872

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cttgacagct ctatggacca tgctatcatt ctctctgcta gcttagtatt tcttagcacc 60
ttcttgatgg ggtgatcctt cttgacgatg atctaagggc tctggaagta cggcttgagg 120
cgttgagcag aggttatgag tgctagcgcc actttctcga tcatttggtta tctctttcca 180
acatcatgaa ggatgtgact gacaaagtag atgggtgttt ggtactttcc atcttcttgg 240
acaaggggtg aactaatggc tttttctgcc actgaaaggt ataggaatag ggatgctcca 300
ngcttangtc aacttataac aggtggtgtt gcaatagtn tctttatagc tagganagct 360
tgcttacagg ctctgtttca caagaacgac tcggttttcc tgagtagctt at 412
```

<210> 873
<211> 408
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 873

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agcttgcttc tacactaaga agcactctat attgagtga tcaccaaaga gagaacaacc 60
accaaaattg aggaccgttt tgtaattntg taatttacia ttactttacc ttcatttctt 120
tcaagttttg taacaaaaag gcctttcatt ggaagtgtgt tgggagcctc caataagtta 180
ccaaacttcc atttgtgtgt aataattcta ggcaattttt ccttaagata gtgagtgttt 240
tgttgggaac cttgaatgtg gtcattccaaa cactcttang atttgcctag ttacatttcc 300
ttgcttactt tcatagctta tttcctttac ctccctttt aaaaccacct agatagtttt 360
ccttttacca attagtnttt ttaccttacc tttcacacct cttttagt 408
```

<210> 874
<211> 321
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 874

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tagccctaga ggngatggac cttntcatgt tntggagagg atcaataaca atgcttatag 60
gttggaacct ccataagagt atggagtcag caccactttt aacatttctg atttaattcc 120
ttttgcaggt gaagctgata ttgatgagga ggaactaaca gatttgaggt caaatccttt 180
```

tcaaggtgga gtggatttta atagcacaca aaagtcaagt caatttaact cctttttaat 240
 agcaaaacaa gtcaattcta catgtaataa tacaatagaa attgtctcta gctaaattaa 300
 aactaagtta atcttgtatc t 321

<210> 875
 <211> 463
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 875

gctttgatca aagtgttcga gggggagcaa gcaaagatag aggataactt tcttcttggg 60
 aagtttgagc tttttgggtt cactacatcg ccaagaggag ttccacaaat caatgtttta 120
 tttgatgttg acgttgatgg catcgtagaa ttcattgcta gagataaaat catgaggatg 180
 aaaaaaagga tcatgatcga caacaagtac tggaggttga gtccctaaga gatgaggaga 240
 atagtgagat atgcaaagag gtataaggca tangatgtgg aggtaagggc aaacggaagg 300
 ccagaacttg cttgagaatt gtgcttttga aatgatggac aaagtgaaga atcttaagaa 360
 attagtaccc atagcaacaa tgttattntt tttagtttca ttaacaattc agtaaaaaaa 420
 aataccgtgc gctaacttga aatgcccncn gcacacatgg ata 463

<210> 876
 <211> 510
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 876

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 gcccctactt ttgacgggaa actccacact tatgaagact atcccgggca agacgatggg 120
 gagggagata cccatcttgg cccctgctc cacctcaaag atccatcccc acatgaacta 180
 cccagccga acatagtccg ccatatcccg gtctcaccca caccgtaaa agaatctgtt 240
 cccttcgcgg aagataaggg aaagattgag gcgcttgaag agaggttaag agcagtcgag 300
 ggccctcggtta attaccatt ctcgatattg gcagaattat gtcttgtgcc caacattgtc 360
 atcccctcca attcaaagta ccaaactttg attagtacca agggatgaca tgttcaaang 420

ggcatctcgg atgtatttgc tgagatggng catattctgc ggacaannag tcgtgggtcat 480
 ttcttttcag acaggttgct tggacngctg 510

<210> 877
 <211> 383
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 877

agcttagccc tagaggggat ggaccttttc atgttttga gaggatcaat aacaatgcct 60
 ataagttggg cctccatagaa gagtatgcag ttagcaccac ttttaacatt tctgatttaa 120
 ttacttttcc aggtggagct aatattgagg aggaggaact aacaaatttg agatcaaata 180
 ctcttcaagg gggaggggat gatgcaatcc tccctaggaa gggccagtca ctagagacat 240
 gagcaagagg ctccaagagg attgggctag agctggtgaa gaaggcccta nggttctcat 300
 gagcctcatg gtagatttct gagcccatgg gacaaggttg ggtctaatta tctttgtaca 360
 tattaaacta ngatgtcatt ata 383

<210> 878
 <211> 490
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 878

ngttgttgtt gtggacctct atnntgaaaa cctccataag agttgttaga aggacctaca 60
 aaagaagtgt tagtgggacc cacatattgt ggtttccagc cataaggctg atttgatctt 120
 gagccacctc tatagccttg gtagcttcct tgaaagtnt gttgaggtct agcttgggtc 180
 tgtagatatt cggcttcctc ttcttgttgc catcaccagg tgctgaacag tggccattct 240
 gatggatacc accacaaaaa tcacatctca naattntttg aacttgggtg gcttgatatg 300
 ttttctatgg tccaccttcg tgatattgct gagtcagttg gcctatctac tttgttaagg 360
 gcctcaattg ttgcgtcaag agnttgtnt gagctagaat ntcactctga gtgtccagct 420
 ccattatacc tattctttga gtccgatent catcatgatg actttgatat cactagctac 480
 tatggagtta 490

<210> 879
 <211> 472
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 879

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 gtggccttaa gcaagctttt attgctagtg gaagtgagga ttcacaggta tgaatcttca 120
 atcatcattg atcaaaatgg gaaaatattc gccagtgtaa ttttttataa tttgcaaatt 180
 gaaagccttg atgtttatat gtctctgtta aatgtgtttt atttgctaag tttttatagc 240
 tgtctcaata atttggttaa ataagttcaa catgcacttg atgcatgcta tcgaggatca 300
 ctaaaatatt ggcataaaag acccatgaaa tgggtccttg tggctctgatn tactggactt 360
 gaatgaattg aactacacat cgctataatg ttcaagagtt cctggcttct gcaatattat 420
 tctagtttat cttgataaaa ctaggaacat ctcgattgat aatgctggaa gt 472

<210> 880
 <211> 284
 <212> DNA
 <213> Glycine max

<400> 880
 cagcttctcc attatctatg ttctcgattg tatctagcaa ccaagttagg gtggagttct 60
 tattatgata ttgtaaagct gagccattgg ataccaattc atccttagct tgcgaggata 120
 agtgcgttaa tctctccatt agaatatgca tatgcgctat cgcgtgatct ctatcaacaa 180
 attcgtctct gtcttcgtag tccatgggtg ccatcaacat accatcaaac atctcgtcta 240
 ggcataaatt gtctatcatc ttttgcgat cctgcctagg atct 284

<210> 881
 <211> 410
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 881

tattccatat tggggaatct aaaataacag acaattgatt gtacaaaaca atgataggag 60

taaaatctag taacatcata atttaaactc actgcaacgg ataatgaca atcaccatac 120
tcaagagcct catggataat gataagatct tcactaccct aatctaagaa aattagacga 180
taaaacaatn tatgtttggt gattacatag tggcactaat atgtaaatgc aaaaatgctg 240
acctggccat gttcattgaa agtatcgagt ccaacaattc caaggtcaag atctccagat 300
aacaattttc ttgtgatgta tttgggcctc ta~~aa~~accaa ctttgagt~~tt~~ ggatagctgc 360
aataatgcaa gagaattggt ttatgg~~tc~~at gtagagatat agtacattga 410

<210> 882
<211> 371
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 882

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acggacatgg ctcccgatcg gaaccagctt cagggtatga ctaaacgaga gcatgagtcc 120
attaaggaat atgccccaaag atggagagat tntgcagctc aagtcgtacc gcccatgatg 180
gagagggaga tgatcacaat tatggtagat acgttaccca cattccacta tgaaaagctg 240
ataggctaca tgccagctaa ctttgcggat ctctgtcttcg ccggcgaaag gattgaatcc 300
gggctacgaa naggcaagtt cgaatatgct gccaatatgg cccccaacaa caagagaaga 360
gccccagtag t 371

<210> 883
<211> 254
<212> DNA
<213> Glycine max
<400> 883

cgcgaaacttt gaccattggt cttccttccc gcaatgcttc ttttcatgtc tgctgagtg 60
ggcttatagc ctaaaccata cttcccacga tttccttgag tatttatcag gctagttatg 120
ccgccgttgt tttttcctaa acccatcccg gggtcaaaac cgttcccaa cataactcgg 180
gccatcatta ccgctgcac ggacagacaa gggtgcccaa agagggagtc cacggaggaa 240
atgctgacca cctc 254

<210> 884
 <211> 101
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 884

ntgagggtgc gtagcccacc atctnttcat agtagagtat cgataatgtg tctaccatca 60
 cgattatcgt ctcccctttcc atcattgggg gtaccacttg g 101

<210> 885
 <211> 300
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 885

tttccatattg tcatcaaaca taaaaagggg aaagggaatg tagtggtctga tgcactgtct 60
 aggagacatg ctttacttgc tatgcttgaa actaagttgg ttggtctcga gtctttgaaa 120
 gacatgtatg tgcattgatg ggactttgct gaaatttttg ctgcatgtga aaagtcttct 180
 gaaaatgggt actataggca taatggattc ttggttaaag caaataaatt gtgtgtgcct 240
 aagtgttcca ttagagagtt gcttgtgagt gaatcacatg agnggggggt gatgggacac 300

<210> 886
 <211> 331
 <212> DNA
 <213> Glycine max

<400> 886

tccattgtcg aatttcgagc gtctcgatat atgatgcgcc taattctgac ttccgagtga 60
 agagttatga ccatttgaat tactggcgag cttccgttga tcaatttcga gcatctccaa 120
 acattatgcg ccttaatcgg acatccgagt gaaaagttat gaccatttga agttctcgag 180
 agcttccggt gttcaatttc gagagtctcg atatattatg tccgtgaatc tgacattcat 240
 gagaaaagtt atgaccactt gaatactcga gagctttcgt tgcgcattt cgagcgcctc 300
 cgtatattat tcgcattaat cggactttct a 331

<210> 887

<211> 233
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 887

aactggggca aataaagagg gtgaggatga gggagaaacc catgctgtga cggccattcc 60
 tatacggcc aatttccac caaacccaac aatgtcatta ctgagtcaat aacaaaccac 120
 ctcttacc accaccagt tatccacaaa ggccatccct aatcaacca caaagcctgt 180
 ctaccgcact tccaatgacg aagaccacct ttagcacaaa ccananaaaa cac 233

<210> 888
 <211> 336
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 888

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 aaaaacatca tttctttctt tttctttctt ccaaagccaa ttctaaagt ccaagcactt 120
 tctccatcac ccacagccac cattagccac cacaaaccat cgttggtctc cattgaaacc 180
 ccacaccgag aggaaccctt caaccgaagt ggaatcttcc aacttggtt gcggtttcgg 240
 tagagaacaa aaccctaata tgacctttcg tttcttttg agactatntt agtctcaaaa 300
 ttatcaagaa ctacgtaggt ctgagttcct catcac 336

<210> 889
 <211> 563
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 889

attgaacca tttaangccc tctattagct gaactctcat acatacgtct gagccggact 60
 ctgactctgc attaccttgt cccangtgat aatgtcaatc cttactcttc gaagcancga 120
 ggaaacgaga gaaggataat ttccactcta aggacataag gagaggaaa gatattcctc 180
 atcaaagagt gggagatagc tatacgacca gatagataat tccaatcca agactgtgag 240
 agagaacaag agaccgagat gacngaagga tagctcctga tcaatgatcg aaagataaca 300

gaagaaatgt gcagagggga tctctggaca gacaatatct atacaaatac agaattgtca 360
 ccaaatgaac acaagagaga aaggaaacca taacctacaa gtggtcttct gccttcgatt 420
 accaaccaaa atactgtgcg tcngtgactt ttgtcgctcg cgtcagacaa naactgaaaa 480
 cgaaaacagc cactactaaaa ctatcaaaag ccataacaac aanagccgat aaccactaa 540
 agagtcacgc cacgggagtc tat 563

<210> 890
 <211> 471
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 890

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 ttaaccttta ttggtgttct ttcatttttc tccatgtatc tcctcacgtg tcttgtgcta 120
 aatgttggtc acatgatttt ttagaatttc gacaattaaa cttgctatag aagctagatt 180
 tgattttcta tggttcaaatt ttcttgttct tggtcttgaa ccataattgt gttgactnta 240
 ngtttctttg agttttgtct tgctatttat ttgtggctga aacttaaacc ataaaattct 300
 taaaaaata ttaatgtata agaaaacctc aaaaatctag agtgacatgt tcacctattg 360
 tagttntgtc ataaaagtca tgtctagtca tgaaacttgc catatatgat tctttatgtn 420
 gngctgaatt ctcatTTTct tggttctttg tctaactcat ttgctcctga g 471

<210> 891
 <211> 461
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 891

tcaacatcag accacttcca ggggtgctgga actacttcac atggacttga tggggcctat 60
 gcaagttgaa agccttgag gaaagaggta tgcctatggt gttgtggatg atttctccag 120
 atttacctga gtcaacttta tcagagagaa atcagacacc tttgaagtat tcaaggagtt 180
 gagtctaaga cttcaaagag aaaaagactg tgtcatcaag agaatcatga gtgaccatgg 240
 cagagagttt gataacagca ggtttactga attctgcaca tctgaaggca tcaatcatga 300

gttctctaca gccattacac cacaacaaaa tggcatagtt gaaagggaca acaggactnt 360
gcaagaagct gctanggtca tgcttcatgc caaagaactt ccctataatc tctgggctga 420
agccatgaac acagcatgct tcatccacaa cagagtcaca c 461

<210> 892
<211> 465
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 892

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taagcaggcg agctcctggc agtcaacaga taaaaggaac aaagaccaca aagcaaggag 120
gcttgtggtg gctggccagc tgtgaatctt gtgtgatata tgggtttttg cctctggtaa 180
tcgattacca aggggtgggta atcaattaca aggcttaaaa atgaagacag gaggctaaga 240
tggtctctgg taatcgatta ccaaaggggt gtaatcgatt accaggcttg aaaacgaggt 300
caggaggcta tgagggcttc tggtaatcga ttaccaaggg ggtgtaatcg attaccaggc 360
ttaaaaatga angcagcang ttgtagaggc ctctggtaat tgattaccag tctgtgtaat 420
cgattacaca gaggaatggg tcaactggtaa tcgattacca cgtat 465

<210> 893
<211> 238
<212> DNA
<213> Glycine max

<400> 893

tttgggctag cccatgttcg atactctaca tagaggtagc gtggaacata ccttgcaaca 60
gtgtgtatac ataggtaaata ataattgagca tgaaattcct agtaaagtgt gaatgattgt 120
cttcctaaat gaatgtatga tagtgtggaa tgcctttttg aatgcaaata tgtgcatgat 180
gtaaatagct atccaatatg catataaata aatatgagtg aaacaataac aatttgta 238

<210> 894
<211> 419
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 894

agcttgtttc cctctaagta cttgattctt ggaaagtgat gtgccatcat tttcttctat 60
tttctagacc ctttttgcac cattttaatt actgattggc ctttaattgtc aattaatcag 120
gcagttttat tatttgggct catttagcta atttgatgtt tttaatctaa ttttaggaat 180
taatgaaaca ttgggcttaa tccggatttt ggttatggac ttgaagaggg caaataaagc 240
agcgcttata ttagttaatt tctaattagg aaattttgca attttatttt atgttggtca 300
gtgtttattt cgttntgggc cagagtattg taatagagcc cagtgacttt gagtgactct 360
ttntaaatag cagccttggg attcgtgcaa ggcattctat tatgctattt tcattattc 419

<210> 895
<211> 402
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 895

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ggtaacccaaa anagatcatt ttccccctgcc attcattgat canatgcttg agcgcttggc 120
aagtatgtct cattacaatn nttttatggt tnttctgggt atttaciaat tcatattgct 180
cctgaggatc aagaaaacac cacattcacc tatccctttg gcatttttgc ctataggagg 240
atgccctntg gcctatgcaa cgccctctgg accttccaac ggtgtatgct tagcattntc 300
aatgattntt tagagagttg catagatgtg tntatggatg attntactgt ttatggatcc 360
tcttntngat gcatgttga tagtctagat agagttctta at 402

<210> 896
<211> 396
<212> DNA
<213> Glycine max

<400> 896

atcctctgag tcacctgagg ctgcagcttc tatccaggct catcttggtg gtgaagctcc 60
ttcttccatg gcttattccc tagtggatgg cgccctctct cacctcttct cctttgtctt 120
ccgctgcac tcctatgggtg aaaatcacca ttaaaggacc tcattgaagc tcaaagatcc 180

agcctccata gaagccccac aagcaagctt ccatcagagg aggagctcac ccctcttgag 240
 ccttcttatt ttgatgcagg tacgcacatg gctaaggagg aggatacctc cacaaaccag 300
 attcatgagc catcttctac acttgtacct gatgatgcca caccatctgc accagcacct 360
 gagtcagagc atcctatctc ttaagattca ccaact 396

<210> 897
 <211> 354
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 897

tgtcatgacc acgctctctg gtgcataatt gatctgtcca aacttaacag ctcatctnt 60
 tcgaacatac tgaataatca gccaggagct tacaagcgta caagtggaaa aattactcaa 120
 ctcttacagt atgttctaag tctgagtaat ggaaatacat attgcttact atttaactac 180
 aacgcttact tgacagatga gcagttcatc atagctatcg gaaggtgaca ttatctctta 240
 ttcttaatta cccttaatt tgtacatgca ttattaaaca accttttaaa acaaaaatac 300
 ttcatcaata ttagctctca agtctaaatt agatgccatg tatcatattt atat 354

<210> 898
 <211> 446
 <212> DNA
 <213> Glycine max
 <400> 898

agcttatgct gcaaacatct acaatagacc ttctcaacct caacaacaaa atcaggcaca 60
 acagaataac tatgacctct ccagcaacag gtacaatccc ggatggagga atcatcccaa 120
 ccttagatgg tcgaatccgt cacaacaaca accttatttt caaatgttg ttggcccaag 180
 cagaccatat gttcctccac cattccagca acaacaacaa caacaacatc ccagaaaca 240
 gcaaacagtt gagggccctc cgcaaccttc ccttgaagaa cttgtgaggc aatgactat 300
 gcaaaacatg cagttttaac aagagaccag agcctccatt cagagcttaa ctaatcagat 360
 gggatagttg gctacacagt taaatcaaca acagtcccag aattctgata gataccttct 420
 aatctgtcag aatccccaaa tgtgag 446

<210> 899
 <211> 360
 <212> DNA
 <213> Glycine max

<400> 899

tgtccgcaaa agatcactaa caacgattct aatgttcgag acctcaattt tctctcacca 60
 agtaaaaatg gatcattcta aggtccaacg ctttataatg aacaccttcc aagtaaaaaa 120
 aatagcttga ttcaccctta aaaagaacta cgtatgtctg atttcctctt cgatggaggg 180
 tacgtagaag caagagccct gcttttgtcg acctcacaaa taaaaaagaa ataaaaagtt 240
 tatgtacaca atttcataca attcaataat taaggctggt gtcctttgag acaaacgtga 300
 gaggtgctaa taccttcttc aaacgtaa atacaactccc aatctggaat attcttcatg 360

<210> 900
 <211> 449
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 900

agctnttaac cttgcttctg tgtatgcaat tttaacaggg gacatatattt atgatgctag 60
 tgaagatttt tctggcactg tatctgttga tgatggtgat gataatatct gcagaatttt 120
 gactgttgaa gatagtttgg gtaccaatgt tggagtatat actgatgata cagaagaaac 180
 aacagacatg ctacatgcac ctctccttc tggaccgaat aagagaagaa aattaatgaa 240
 ttcttttagt gctggagttg aagttgatag ctactcgaca gctgaaattg ataactcatt 300
 ggattattct cagacctcta gctgtgtttc tgatgataca gttgaaacca ctcaagatga 360
 tacactngaa accactcaag atgatatagt tgaaactact caagatgata cagttgaagc 420
 aacacaatat agtgatgggc tactgtcat 449

<210> 901
 <211> 399
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 901

ntgagggtgc gtagcccacc atctnttcat agtagagtat cgataatgtg tctaccatca 60

cgattatcgt ctccctttcc atcattgngg gtaccacttg ngccgccaga tccctcaacc 120
 ttttgggcgt gttctttgaa agatccgtcc ccctttntgc aaatgttctg tagttgcac 180
 ctatccagaa ccatatcaaa attgtactaa tactgcctaa caaaggcaac caataggtcc 240
 ttccaagaat ggactcggga aggttccaag ttagtgtacc gggtaacagc taccacagta 300
 agactttctt ggaaggaatg tattagcaat tctcatctt ttgcgtatcc ccccatcttc 360
 tgacaataca tcttttagatg gttcttgga caagtagtc 399

<210> 902
 <211> 565
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 902

attgacgcat ctttangcat cactatcgat actcaagctn gcctanagag gtccaggana 60
 gacaaggcgg ccgaaggaac tagttccgcc ccgtagtacg acagtcaccg ctttaggagc 120
 gttgtacacc agcagcggtt cgaagccatc aagggatggt cgtttctccg agagcgacgc 180
 gtncagctca nggaggacga gtatanctga ttccaggagg aaatagggcg ccggcggtgg 240
 gcaccactgg ttactcccat ggccaagttt gatccagaaa tagtccttga gttntacgcc 300
 aatgcttggc caacagagga gggcgtgcgt gacatgagat cctgngttag gggtcagtgg 360
 atcccgttcg atgccgacgc tatcagccag ctctgggat atccgatggt attggaagag 420
 ggccaggaat gcgagtatgg ccagaggagg aaccggctctg atgggttcga tgaggaggcc 480
 atcgcccagc tgctatgtat anccngntan ngatttgcen gactgctgan ggagagagtg 540
 cgatcatcnc accatatgac acccg 565

<210> 903
 <211> 381
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 903

taggggggag aagtgaagaa gaaaaggggt cagcctctaa ggcacttctc tctttctcga 60
 aattgctgag gaaaattatt ttcgtgaaga aaatccaagc cgaggcgctt ccgtaacggt 120

tctgtaacgt ttccatgagt aattacgga agattctcga cgtttcttca agattcatcg 180
 ttcgttctgc gttttcttca gtcttcaacg ggtaagtacc tcaaaccaag cttttcaatt 240
 cattctatgt acccggtggg gtccacattt tgtttcatgt atttttattc tcattttcat 300
 ttactttnta taccctcttt tgacgtgctt aagccattta tttaagtcatt ttctcgctta 360
 atgaaanaat aaaataaatac t 381

<210> 904
 <211> 387
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 904

cccacacat gtggtactat gtggtgggctg ggcgatgggt cataatgatt ctccacatcc 60
 acaaatacag tataaccacac catccctctg tgcccacctc caactgagct cacgtactcc 120
 cacgtagccc ttatcctcgn tccctctcaac gccgggtccc catcaatcct cccaagcttc 180
 cacaacatcc aagtaattca acattcaatc atcacaaact aacacagcca agaaaaatagg 240
 gccaaaggcag aaaactctgg ccaaaacaca aaccaacatc acagcttttc acattcaatt 300
 acctcaataa gagtctctgt gttccaggtc ggtaaccgct ggatcgaact cgaaattata 360
 ctgggaagct ctagtacata agtctac 387

<210> 905
 <211> 130
 <212> DNA
 <213> Glycine max
 <400> 905

gagccttggt tccctttcct tgttttgaag ctactacaa gccttaaagt aaaaaccatg 60
 atatcaccat atccttaagg aattttggag ctttggaatt gttttgggaa taagtgtggg 120
 gggggggggggg 130

<210> 906
 <211> 417
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
<400> 906

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ntccttcaca ttcagcanat tcagcattta aatgtgatat ttaatgttat gctnttttat 120
gccatgataa ttggtggaat gaatattatt tatttgtcaa ggtttcatga tatcgaatat 180
tgatacctaa naagggtaat atttcaagtt gtgtgattag tggtattttg agatgaaaca 240
ccaactatat gtaatcttat ctttgcatta tcaagttggg attaaaaatt tgtaatctat 300
tcgttgata tgatagtagt agggactcat aaggatntac ttagtaagag gcttaacct 360
aagtaagaat ttgtttttct gagacaaaac tgcagagatc atcntgtttt attatta 417

<210> 907
<211> 465
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 907

cgtgcattca atatcctgat gaggagggtc catatgtctc aagactggac taatacattt 60
gctgtccaag tttcatgggc ttgcagggtga agatcctcat aagcatctta agaagttcca 120
tattgtctgt tccaccatga agccccctga tgtccacgaa gatcatatct ttctaaaggc 180
ttttctcat tctctggagg gagtggcaaa agattggctg tactaccttg ctcccaggtc 240
cattaccagc tgggatgacc ttaagagggt gttcttgggg aaattcttcc ctacatctag 300
gaccactgcc atcaganaag acatttcagg catcangcaa cttagtggag agagcttgta 360
tgagtattgn gaaagattca agaaattgtg tgcaagttgt ccccaccacc agaattttga 420
gcaactcttt ctgcaatatt tctatgangg acttancaac atgga 465

<210> 908
<211> 588
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 908

gggaccacgg ggnnnnaatt gatcgcatgc tattangcca tactatagat tactcaagct 60
cgagttgagg aagtgtagaa gggtgaaact tcttggtttt attcgttgac cacagagtg 120

tacctggaga tatgtcgca gggtcaggag aaccttggga cgtcagggtg tgtgctattg 180
 cccaaaacca agcttgacca atcccgaccc aaccgggca tagtcgggtca gtgagaacct 240
 gtgatgtacc taaacaggcg agctcctggc agtcaacaga taaaaggaac aaagaccaca 300
 tagcaaggag gcttgtgggtg gctggccagt tgtgaacttt cattgatatg tgggttatgg 360
 cctctggtaa tcgattacca aggggtgggtg atcgattaca aggcttaaaa atgaagacaa 420
 gaggctaaga tggctctctgn gtaatcgata ccacggngtg taatcgatta ccaggcttga 480
 naacgaggtc aggaagccat gagggcgctct ggtaaatcga taccaagggg tgtaatcgat 540
 taccaggctt ananaggggg atggacattg tganggctct gtaatcan 588

<210> 909
 <211> 267
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 909

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 gatggtcgtt tctccgggag cgacgcgtcc agtcaggga cgacgagtat actgatttcc 120
 aggaggaaat aaggcgccga cgggtgggcat cactgggttac tcccatggcc aagtttgatc 180
 cagaaatagt ccttgagttt tatgccaatg cttggccaac agaggagggc gtgcgtgaca 240
 tgagatccta ngtaaggggt cagtga 267

<210> 910
 <211> 361
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 910

tacggatgga atacttactt ggtgtgatga acaagagcgc gatacagaat ctataaatgt 60
 gcaaaatgat gaccctangg ctgctaactc gtaaataccg tgggtatggc tcttgaaagg 120
 cgaaaaaaga agtttatgaa tgcaaaaacg cgcccccttt cgtcattctt atatattggt 180
 gcanggggtgg ctgcccagg cgagctaacc tgcattatct tttttgagag gaacattaac 240
 catgtccact ccttcctttt agcgctttgc ctaacttgaa cttacttaag ttagaatcaa 300

gcgttgatta cttatTTTTta ataacaaaca gatagtaaga taactgCGaa tacaaaggat 360

a 361

<210> 911

<211> 471

<212> DNA

<213> Glycine max

<400> 911

agcttctcct actgcaattg tcaaacacgg tgtctaaatg tgagttCGaa taacaacttc 60

ttttacatgg tgatCGcgga tgtacatccc aacttttagtc atattcagat tctCGttcaa 120

actagCGtcc accattgcac tccaacctac tcaatatagg agggctccat atttcaattg 180

tgctataacc acgcattctc tctaattctcc ccttgcacct tttttcattg aaccaatcct 240

agaaaaata tttgcataat gaatcacctc cgaatctgta ctatccttat tgctccaaag 300

cttttcattc catctcctcc aaacactcca taacatcata gcaacacgtt tcccttgccg 360

gactgatagg acttgaatta atctgaatat gaattcaaag cacgaactct catcaaatat 420

cgttaggtcc tcaacaacat cccaagtgga agactcttaa gttaaattatt t 471

<210> 912

<211> 409

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 912

tatgcattgt ggaatttcac cagagagagc attgtttgaa acatctcaac ttgttgatgc 60

tttcagctta accgacttca aagggttaggc tttccattga acgagttgtg ggacaagttc 120

aataacaatg acaagtanga aagaccaatg acttgcaaag gaatagttcc actaattctg 180

ttgtgtgaaa ggtcaagggc attgttaaatt tggcaattcc caacaccagg atgtatgctt 240

ccttccaaca cattatttga caagtcaagt tgaaacaaaa gactaagggt gccaatggat 300

aatggaattt ctctgacag tttgctcaca tttaaattca atgactgcat cttttggaac 360

atgacaaaag aagcaggaat agtcccagta atctgatggc caccaatat 409

<210> 913

<211> 442
 <212> DNA
 <213> Glycine max

<400> 913

gatcttaagc acctgcggt gcagctttgc ttaagacatt gtcttggttg tttgcttctt 60
 tatttttttc tggaaattgc tagtttagta taggtccttg atttttggtt tatttgtaat 120
 aaatgtgtac tccttggtt tgaggcttaa agcttaagta tagagtagtt gctttcaaga 180
 atagtgttgc tatggaaatt tcctttaaat ttgcgggcaa cgtcaaacca aaatcctacc 240
 caatgttttg aaatccatca tactgcgcct ttagaattcg aagaatggta caatgatttt 300
 aatgggtccc accactgggt tatactgtat taaatatcca ataaatatac ataaataatg 360
 gaatcacgtc ttagacagaa atgttacata acaatactac aaataatcac atactactat 420
 gctacgaata atcactgatc tg 442

<210> 914
 <211> 524
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 914

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 acctggagat atgtcgcggn ggtcaggaga acctggggac gtcaggtggn gtgctattgc 120
 ccaaaaccaa gcttgaccaa tcccaaccca acccgggcat agtcggtcag tgagaacctg 180
 tgatgtacct aaacaggcga gctcctggca gtcaacagat aaaaggaaca aagaccacaa 240
 agcanagagg cttgtggtgg ctggccagct gtgaactntg attgatatgt gggttatggc 300
 ctctggtaat caattaccaa ggggtgggcaa tcgattacaa ggcttataaa tgaagacagg 360
 aggctaagat ggtctctggt aatcgattac cacgngtgt aatcgaatac caggcttgaa 420
 nacgaggtca ggaagctaag gaagcctctt gtaatcgatt accaaggggt gtaatcgatt 480
 accaggctta naaaggaac tgggagatga tggaagcctc tggg 524

<210> 915
 <211> 305
 <212> DNA
 <213> Glycine max

<400> 915

ggctgcagct tattgctaca aggcacttac tcttctagcc ccaagagact cagcataagg 60
atgcacagac caaagttgtg ttgtataaaa aatgtgttga ccaatggaag gtgctaattg 120
caaaaacaaa tgaaagctat gccaaagcaag ccaaaaaaaa aaggaaggaa gtggttcttg 180
aaccgggaga tgatcttggg cattcgagga caaatgtttt ccaagaggga gggaatgatg 240
agaatcatga aacaggccaa atacagtcta aaggcccaag tggagaagga cgaaggccca 300
agtgg 305

<210> 916

<211> 353

<212> DNA

<213> Glycine max

<400> 916

tcattgccta acaagccaac ttacaacatc tagccccaag agactcatca taatgatgca 60
caggtcaaag ttgagtatga gataagattg tatgaccaa tgaagggtgca tattgcaaag 120
aacaatgata gctatgccta gcaagccaac aagataagga atgaagtggg acttgaaccc 180
tgtgatgac ctggacattt gaggacaaat gttttccaag aaggaggga tgatgagaat 240
catgaaactg gccaaataca cgctaaaggc ccaagtggag aaagactaat gcctgagtgg 300
agaatgacaa taaccctgag tggagaatga tgaaagccca agtggagaat gat 353

<210> 917

<211> 404

<212> DNA

<213> Glycine max

<400> 917

actcagcttg tcatgaccgg tctcttttgg gcattattga tctgtccaaa cttaacagct 60
cattcttttt gaacatactg aataatcagc caggagctta caagcgtaca agtggaaaaa 120
ttactcaact cttaaagtat gttctaagtc tgagtaatgg aaatacatat tgcttagtat 180
ttaactacaa tgtttacttg acggatgagc agttcatcat agctattgaa aggtaacatt 240
ttctcttatt cttaattacc ccttaatttg tacatgcatt attaaacaac cttttataac 300
aaaaatactt catcaatatt agttctcaag tctacattaa atgccatgta taaatattat 360

ataaaaagttg ttttcatatg ggattgataa gcgtgtgtgt gtct

404

<210> 918
<211> 453
<212> DNA
<213> Glycine max

<400> 918

agcttggcgg caaccacctc cctttttttc tctataatag gggaaaaagg gcagagtaat 60
ttggctcaac ctttctggaa tttaggattc tcttgaaatt agagagaaaa attgtttccg 120
tgaagaaaat caataccgac gcccttccgt aatgcttctg agacattttc gtgagcgatt 180
ttgtaaagat tcttcaccgt tcttcacgcg tcttcgttcg ttcttcgtcg ctcttcggtc 240
ttcaaccggt aagttcctga aataaaacct ttcaattcat tctatgtgcc catagtggtc 300
cccacctgtt tcacgtgctt ttattttcat ttcgtttctg ttccgtaccc ctttttgacg 360
tgctttaacc attatttaag tcgctttctc acctaatcaa gtaataaaat gagattccac 420
caatcatttg agttgtaata tcgtttaatc tct 453

<210> 919
<211> 419
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 919

gcngcaccgc tnaaggtaaa gactcttcct gtggccttca aacttccaat atggtcattc 60
aggccctcac cattctgctc cttcttggga tatgggcaat ctctctaaat atgccccctc 120
tgtcaatagt tgatacaagt cgcgccttta tcagcataat tcgaggaaat gtgccttggc 180
ttaccacatt tgtaacaagt gatctgagtt gataaagaag tgggtttgct accattacca 240
ccagcaaacc ccatagcatc agtcctctga ttgttggggc gattaccata tgtcttaaga 300
ggggttgagt acgatcttcc ccgttgggtga ggtccattct ttttgttctt cattngncct 360
gcactcctat aatacgtgc cttgtatcag aagcttcac ccaatccgga catgttacc 419

<210> 920
<211> 245
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 920

cacaacaagc tttcacatcc acaatgcgcg cataaaccca ccatcccctg gtgcccacct 60

ccaactgagc tcacgtactc ccacgtagcc catatcctcg tttctctcaa caccgggtcc 120

ccatcaatcc tctcaagctt ccacaacatc caagcaaac aacattcana cagcacaagc 180

tatcacagcc aagcaaaaca gagcaaagc agaaaactct gctcaacaca tcaaccagaa 240

tcaca 245

<210> 921

<211> 397

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 921

cgctaatta acctgatatt gagaganaat gattattaaa tacacaaaat ggaagtacta 60

agtatttatt atctatattt aatagaaaat acttataaca ttacaaaata accataaatt 120

ggaagagttt gatacaattt acacaagttt tatacacaaa agttcgtcgt attcaccgac 180

taacatagca caagacatat ccgtggaggg ttctgagggg atagtcaata acatcacgac 240

caacaattac ctcactttcg ctgacaaaga gatactcgtc gagggcaggg gacacaatgc 300

acgtgtctgt caaatgtttg gacaacatan gggccaaagt gctcatcgac aatggctctt 360

ccctcaatgt catgcncaaa gctactttgg acaagct 397

<210> 922

<211> 406

<212> DNA

<213> Glycine max

<400> 922

ggctgcagct tctcgatata ttatgcgcc aatcggacc tcagtgtgat aagttatgac 60

cattttgaat tttcgagagc ttccattgtt caatttcaag cttctcgata aattatacgt 120

ctgaatcgga ctttcgtgtg ataagttatg accatttgaa ttctctgaga gcttccattg 180

ttcaatttca aacttctcga tatattatac gtctgaatcg gactttcgtg tgataagtta 240

tgaccatttg aattttctga gagcatccat tgtttaattt caagcttccc gatataattat 300
gcacatgcat cagactactg tgtgaaatgt tatgaccatt ttaatttctc gagagcttcc 360
gttgttcaat ttcgagcgtc tcgatatatt atgcgcctga atcgga 406

<210> 923
<211> 393
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 923

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nattgtctta tactttcaca cggaacaccg attcaagctc ataatatatc gagactctcg 120
aaattgaaca acgaaagctc tcgagaaatt caaatggtga aaacttttca gacgaaagtc 180
ggattcagac gcataatata tcgagaagct tgaaattgat caacggaagc tctcgagaaa 240
ttcaaattgt cataacttgt cacacggaag tccgattcag gcgcataata tatagagacg 300
ctggaaattg aacaacgaaa gctctcgaca aattcaaattg gtcataacta ttcacacgga 360
agtctgattc aggcgcatac tatatcgaga ctc 393

<210> 924
<211> 382
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 924

atgccccaca ttattttccat gacacaaatg ccaaaatgat gatttggaaa cttcatgcaa 60
aacttgtcat gcatgcatct atgcggacac tcaaattgtca aatttttatg gtcattgtgat 120
gctaaggctc aggattcatt tctctatatt ttaatcaacc caatgtttcc aaaatatggt 180
cttttatcaa tttgtgcatt catccgagtc catttcgggc gtccgggaaa tttcacagca 240
ttcacccttc aggcgtagac acatttccca aaaattggtt atggatcaatg aatnttttca 300
aagaaaagtt ggaaatcgtc tcttttcaaa agcatgtcat ttttagctag acaacttatt 360
ttctttnttt ctctttcttc tt 382

<210> 925

<211> 386
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 925

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 tgataccttt gcgagattcc tcacggaaaa cgttacggaa acgtttcgga agtgcctcgg 120
 cttagatntt cttcacggaa acaattnttc caagcaaatt cgaaggagag agaagtgcct 180
 aaggggctgg acccctttct tcttcatttc ctcccctatt tatagcaaaa taggggaggt 240
 ggttgccgcc cagctcgccc aggcgagctc agctcgccca ggcgagcagg gttgcttctt 300
 ccagaagcaa ccgccttctg gaggaatatt ccagagggcc caagtgggcc tgggtgctat 360
 ttgcaccnn cattttacta agtaca 386

<210> 926
 <211> 419
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 926

cgccatnntt tgtttaatat gtntnntttt tcaataaatg gaacttcatt ttatacataa 60
 ttggtattag tataactatc aagtttcaac tattagaaat taaactagac attataactt 120
 ttaaagcagt tactattata agaattaata ttttttcata atatatagca atccatgatt 180
 agtttacagt atacaaaata tntatttcat taatatattt caattaaatt cttgataaat 240
 aaagacacan ttttaacatg atctatcgtg tatatgaaag tgtcttcggg cagaatataa 300
 ctctaacaaa atttctaaaa catagatata tacaatatca tataaaatat aaaataataa 360
 aattttaaac tacaatggca aggtntataa tgttataact tcgggtaaca aaaaaaaaaa 419

<210> 927
 <211> 438
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 927

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ggcgtagctg cttactgtga aggcgccgaa actgtgatgc cacaaacttc aagtcacgcg 120
 ctttcagggc aaaggcttcc acttctgaaa gggtttggac agtcctgggt gaggtangaa 180
 ggttgggtga agaagtagga tccaaagccc acgtgagaag ctctctcca cagaagtcac 240
 cagccttgag gtactcagag ttgaagaagc cggttcttcc accgttagtt gtcattggta 300
 atagcttgcc acgcattatg aagagcatct catcaaccgg atctccctcc cggacaatgt 360
 agctttcttc tgtgtaagca ctggcttgag aaagcgcaca ttgatncaga agtggtcgtc 420
 atttctcaac attggacn 438

<210> 928
 <211> 491
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 928

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 antttgaatt atggccttgg tttgtgtgtc tagattgaca ccagactcct aaatacacia 120
 tatctttcat tgcaagctta gcaactgtcc caaaacccaa gtttattcga aaccaagtgt 180
 catgatttct atattaccaa ttntgctagt tgtaaatgtt gaatcatagt tttgctctct 240
 catctgcctt ttgtctcatc tctttacctt acaacttagt caattctatc attacccttt 300
 ttcaatatgc agaatcagca acatgcaaac atatctaac cagcaaagtc caccatcaat 360
 agccaggcta tggccagaa ccaacaaaat gctcatgtc ccatttcttt catcttctaa 420
 atntattgga gcttctgcag attaaaagaa gcattngttc ttcatttcac atgaatctac 480
 tgggttagtt a 491

<210> 929
 <211> 348
 <212> DNA
 <213> Glycine max
 <400> 929

agaagaagtt catagagatt gattggattg tcagaaagat tgaattgatt gaaaatgcaa 60
 aacaaagcct tgcttttata gactcttcgt gtctgggtcaa gaagaccact tagaagagtt 120

ataactttta gaaaaactta aaaccaatth gaaaaagtca aaaccttttt gaagagttac 180
 atcttttgat ttattcagaa acaaactg gtaatcgatt accaaattag tgtaatcgat 240
 tacacaaagc ttttgtgtga aaggatgtga ctcttcacat ttgaatttga atttcaacgt 300
 tcaaaggcac tggtaatcga ttaccaaacc attataatca attacaac 348

<210> 930
 <211> 408
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 930

ctgatcanat gactaanatt aatcggcaca acagatattt agcagtagga gggaggcctc 60
 caacatacac ccgcctagca tatcatgtac cctgttaaatt caatacataa agacaggcca 120
 taatcaggga aagtcagtct aaacttaaaa tgaaacttat ataagactgt acttagtctc 180
 ggcattgcctc cactactcgga aagactaaac ttaaaacgtt ttgtgttgag gaagttagt 240
 tgtctctact ctgtatgtat aatgactctn ttcttctcaa tgaaagagaa tatcttcttc 300
 cagtagcaca atgatactat acaaacaaga gcgcaatata nagaaacaca tggtaaaaga 360
 agaccacaca cctttcatct acagcagcac atgatactgt aatatatg 408

<210> 931
 <211> 455
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 931

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 gatcaccttg accaatggaa tctccttccc tcttaggtgc tttgttcgcc tatctttgat 120
 cctcaaaggc aatgttccat atgtcaagtt cttcttcact tgtacgtcat ccaatttgat 180
 cacacgagat ggatcatgga tatactcacg aagttgagac acatgaaaga caatgtgaag 240
 gttagaaaga gacaggggta atgcaatttg gtatgccac agtaccgact ttttttagaa 300
 tttggaaagg acagataaaa tgaggtatga gttattgnga tttcaatgct cgaccaactc 360
 cagtccacaa agtgactctc aagaatacat gatcactaac ctcgaactcc aagtctttcc 420

tcctttcttgt cctgatagct ttctacctac tetga

455

<210> 932
<211> 328
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 932

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agtttgggtg gtaattacta tggcttagtt atagtagatg attactcaag gctcatatgg 120

actttgtttt aacaaaaaat gaagcttttg gtggctttta aaaacttgcc aaggtgattc 180

ataatgaaca aggtctcaac attgtttcac ttagaagtga tcatagaggt gaatntcaaa 240

atgagtcttt tgaaaactnt tgtggagaaa atggaattca ccataatttn tcttgcccaa 300

gaacacccca acagaatggg tttgtgga 328

<210> 933
<211> 445
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 933

agcttggaga tgatgcttca atggaggata agtaagagag aaggggggag cacgaaattg 60

aaggaataaa agaggggagag aagttgaact ttgaagtgtg ttcataaga ctttcattca 120

tcaaagttac aacaagtgtt acacatgctt ctatttatag actaggtaac ttccttgaga 180

aaacttcctt gagaagcttc tttgagaaaa cttccttgac aagcttgagc ttagctacac 240

acactcctct aataactaag ctacacctct tgagaagctt ccttgagaag attcctaaag 300

aagctagagc ttagctacac acaccncta taatagctaa gtcacccca tgccaaaata 360

catganaata taaaaaaaag ttcctattac aaagactact canaatatcc tgaaatacaa 420

gggtaaaacc ctatactact agaata 445

<210> 934
<211> 458
<212> DNA
<213> Glycine max

$\frac{d^2}{dt^2} = \frac{d^2}{dx^2} + \frac{d^2}{dy^2}$

<210>	935
<211>	401
<212>	DNA
<213>	Glycine max

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ttgttcttttc tttgtctaac atacatacta gctcaaactc atgaaaagaa acacaaactc 120
catcaaaatc atgcactcaa tttaagtact tgtagttttt cgtgagggaa aatacttgta 180
cttggggggca tgtcactcgg tttggaactc ccttgtgact cgggcttata accattgggg 240
gtgggggtgga gttgcctgtg cacaacagga tgaccttgac acttgcctacc cagctttctt 300
gggtgtgagt gtcgtgtggg aatgctcang ctatttcattg acgaatggta ctacattgca 360
tttgagagtt aaggtcaagt gcatgcatca tactaagcat g 401

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<223>      unsure at all n locations
<400>      936
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384

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 tccagaaaag gatgagaatg gaggattgcc ttgagggtcc tctcttango aatcatgaaa 240
 cacaactcca aactcaaaag cggaggacac atgaacagcc ctaagcaata acattcatgt 300
 ggctccggan aaggacaaga atggaggaat gccttgaggg tcttctctta agcaatcatg 360
 gaacacagct ccagactcga aaatggagga cacatgaaca gccctaaagc ataacattca 420
 tg 422

<210> 937
 <211> 506
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 937

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 tagtattaat gtatgtaatg tataatntag tggngaatat taaggctata ttaatgatga 120
 tatangattt cattagaatt agaaaaaggg gtaattaacg tcatatagag tctaaaagtg 180
 gagggcattt ttggtaatga ctatacaact agtttaaaaa taggatttta atttaattaa 240
 ttgggtgacta attaaagtgt ctaattatta tgatgtaaat aattaanata agttagagtt 300
 gaacaccctg^gaanattataa ctcagactga cataaaactc tatgtngggc atctgtgtgt 360
 gtatgaagtt aatttcagta gctataccgt tttaatcata gaantntcgt gctatgatat 420
 atgtatgtga ctggtttagt aagcttgact gngaataga ctacctttgc tagattcatc 480
 agtgcacatt tgactgtgat taagcn 506

<210> 938
 <211> 397
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 938

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 acgagacatc ttgccaaaca aagtcaggtt agcgataact cacctgtgct ttttcttcca 120
 tgctatatgt agcaaagtca ttgatcctgt caagtttgat gagttggaaa atgaggccgc 180

aattatacta tgccagttgg agatgtatTT tccccctgct ttctttgaca tcatgattca 240
 cttgattgtg catctggTca gagaaatcaa atgttTggT cctgTttatc tatggTggat 300
 gtacccgatt gagcgataaa ttgcanaaga agccattgaa tttttttcag aataacttaga 360
 gaatngctaa acctgtggcc ttctgagtct cgcattga 397

<210> 939
 <211> 422
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 939

agcttgtaac ttatataata tatacatTTT attgtaatta tattttaaca catcagaatg 60
 gtgcgcccatt aaccacagg tcccaggatc gaaacctggT tctgataaag agtggcttcc 120
 gatctatcac atatatatat attttatgcg taaaacatat atcattacgc aatgacattt 180
 gagtataata aaaaatagtt ctgcagggcc taacatttca gtgcttatat taatttagtt 240
 accatttaaa ttttattatt gagtcaactt tttaacgtat attcatattt tctctttggT 300
 aattntattt taatttgctt aagtaaacad attttttatg gataataatg gcttccagtt 360
 tcttagtgaa ccacatctga aaaattatac ttgaacaaga agatgtgttc actatgtcat 420
 ag 422

<210> 940
 <211> 427
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 940

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 gcaacgtgta agcattgcgt tagctataat ttatgaagaa ccacttctag ttctataatt 120
 gtacaacata ttagcatatg ccaaactatg tgtatcattt ggatcaccaa aataagaata 180
 ttacctcaat aaaatctcct ttTggcatta gtgctctgca tgcattcteta ntcctttggT 240
 atggTgatat taaactagtg atgcaaataa caccagcatc tgcaaagagt ttagccacct 300
 cacctganaa ttntatatcg tgatgtctaa ttattaataa aacaataaaa tataatcgga 360

agatatcagg gaaagcattt agaaagcaac ataagaaaaa acagataaac tcaccaatcc 420
 ttctaata 427

<210> 941
 <211> 119
 <212> DNA
 <213> Glycine max

<400> 941

tgcttgagag acttctatgg atgttggatc tttgagcttc actatatgtc cttcaatgg 60
 gattttcaat catggagttg catcggaaga taaaggagaa gaggcgagag gaggtgtca 119

<210> 942
 <211> 473
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 942

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 gaattacctc ctcgggttgc aattgcactt agcgcacagg tctcggttaag aaagttgtcc 120
 aaagatgtta tttgaaaaat ctcaatagta aaaaatgtag gcatgaatca agaaagttgc 180
 agttcatggt tgaaggtgat ccaacggtta acgagtctgg gatcatggtt ttactgaaat 240
 aggttaaaca aactccacat aaccttattg ttcacaccaa gcaaccgcac acaaataagt 300
 cacacaacac ctcaactaat ccaacttaat caaagaatgc aagaattata ttaaaccatct 360
 attttcagtt atcaatattn taggctgtta caaaagacct tttcttgggt atcaacacca 420
 aagtattcaa gaatcttgga gatcagaact gcatagggac acctganaaa aac 473

<210> 943
 <211> 357
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 943

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 cngaagagaa tntcaaaatg actctgaatt gttttgtgaa caaaatggca ttaatcgtaa 120

cttttccgct ccaagaacac cacaacaaaa tgggattgtg gggaggaaaa aaaagtcctt 180
 tgaggaactt gttagtgttt agctctactg agctntaaaa gattggctaa gatcttgtaa 240
 aaacataagc acttagacaa tgaatgaaag ctggagttgc tgcacatgat gtccaacgct 300
 atgtcaagga ataagatncg gctgcacaat gcacaaggca agataaaatg tcaaattg 357

<210> 944
 <211> 437
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 944

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 tatcaccttc ttaattgtac acatggggca cttgcccccc aaatgcgca agtagaagag 120
 ataattttcc gggctctcgt gtcgtaaaaa tgcattcata tcatgcatcg cataagcatc 180
 tcttcataac atcataatgg acatatcctg catttgctcg ttatcatatt ccagcctcac 240
 attttgcatg agtcatggca tcatcatgca tatgcgttca acaaactttn tgatctgcaa 300
 aattgcatac catttggttt catgtttgct catccttgcg ttttcctcta caaaacaaaa 360
 acaaagaagg gggaagcggtg aaacttcaca ctacattctt agtttcatgt gttaggagcc 420
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<210> 945
 <211> 349
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 945

ctaacctcat catctctcat agtctntaga tntgngagcc aatccaatcc ttgcgtccag 60
 actctcagcc acttatgata gccgccgatg ctcccattac tgcttcccct aagctctatg 120
 tcctttcttc acgccgcac ccatgccttg cgaactcctt ggagtaccct cgcgttggtg 180
 tcaactgaaac cccgtgcat gaaaggcggtg atgctttcgt ctgatggcac tcctctcatg 240
 gggtagccaa gctgtcttat ggcgaggacg ggattataat taatacaacc ccttgttcca 300
 tcaagggaac atttggacat ccttcgcatg aagatagaat cctgattct 349

<210> 946
 <211> 156
 <212> DNA
 <213> Glycine max

<400> 946

cattctctct cattatcata ttagcattgt aggggggttc agagcattta tacttcttct 60
 gtatctcgag gaatggtcta caaaccttga agcctagcgt tagttgtctt atacgactaa 120
 attctgtata gaaaaacctt tgtcacagca tgtata 156

<210> 947
 <211> 387
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 947

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 tagtgtagaa gtaacttata gaaaagagat agttatatag aagagtaaag aaaacactaa 120
 ggttttatac cggttcacct caactcttgg gctatgtcca attgtctttc aaaccttgaa 180
 gggttccatt aatcaattct ttgattacaa tcaggtattc tctatgtcac ttctggctat 240
 aatgagtact ttgtaccact catgggaeta cccttaatct cctcatgagt taagacttaa 300
 gtattctttg tcactaagtc attcctagcc ttcacaaaca atatatgttt gatagaaaat 360
 gattctaatac actcanagag tgttaca 387

<210> 948
 <211> 400
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 948

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 ctacaaaagc ctccataaca taacatcatc ctcaattctc aaaccctaaa caacaaagca 120
 cccccgccac caaacacgca cacacaagca gctaaatatac attcaagaag taggatattg 180
 ccataatcca caaacctaag aaaccacaaa actaaaccaa gcaccaacac cattccaaaa 240

acacctcaca aaacattcac acacacacaa caaaccaaat aaaaaacaca cccaccacca 300
 attcacatcc aaatcccaat caaacaccaa tcaataacac caaatcaac ttccaacaaa 360
 tccaagccaa canacagcca tcaactacta canaccaact 400

<210> 949
 <211> 384
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 949

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 cctagtggat gacgcctcct ctacactctt ctctttgtc ttccgctgca tctccatggg 120
 ggaaaatcac cattaaagga catcattgaa gctcaaagat ccacccctcca tagaagcccc 180
 acaagcaagc ttccatcaag tggatcaga gcacaagagc ttcaagtagg tgctccttaa 240
 acttccatta attnttttgc ttacacttct ctccattgt tggttcttca ttntttctcc 300
 atgtatctcc tcacatgtct tgtgctaaat gttgtaaca tgattcttta gagtttccac 360
 caattaaact tgctatagaa gcta 384

<210> 950
 <211> 341
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 950

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 gtacttttaa gctaatagtt gaagacactc attgtgattg tcgttgcgct ttatctcctt 180
 tatgtttaat tactcatttg accctatag ttatagaaac tttctctttt agtccctata 240
 cttaaaaaaca tccctntta gtcctacac attccatttt tattcccttt cagtctctac 300
 acatcattnt aatcccttgt agccgctatg gtgaggacta a 341

<210> 951
 <211> 464

<212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 951

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 tactactggt ttgcatcaat tttgtcttct gtgacccct tataatgtgg attaattcaa 180
 ttgagcaaat ctgttcgtn tatgctcagc ttctataagt cttttcattt acagatatat 240
 tgtaatatg ttattgagtt cttggtttta aactagttat tgttattgta tctacccttg 300
 atttaaaact agttaaatat ataagatact ataattcaaa agggcagata gatccaaact 360
 tgatacacat tttcacttga gagaatccna gtctgtgtgt aatcatcgcc ttccatttca 420
 atgggattag gtgatggtt tactcatggt atgactctat cata 464

<210> 952
 <211> 309
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 952

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 ccaaatttgg gggaggtatt atcaacgtaa atntgttcca aatttgggga aggcactcgg 180
 taacgattga aatggtcaaa gaaaatagta tatacacact ggctctatta tctgtgttaa 240
 aaaaaaacca ataaaaaact gtacgtataa ataaagttaa taagtgtgta tgctataaat 300
 tcaggcatg 309

<210> 953
 <211> 481
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 953

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attttaggct taaatagggtg gctttgttcg tgcttgagcg cttagecgaa ctctgaaccg 120
 cttagcacga attagtgaat ttcggcttaa cgcgtgcttt tctcgctcag cggatggact 180
 aaagcgggtgc gcttaacgag atgacccttt gctcagtga catgcacagc tcctccttct 240
 tctagattct tctcgcgct cagcggatag ctcgctaagc cagtagattg gcttagcgag 300
 aaggtgaaaa tcagcacctc acaaactttc ctaattaacc tgaaattgag agaaaatgat 360
 tattaacac acaaaatgga agtactaagt atttattacc tatctttaac aaaaagtaat 420
 tacaacactt acaaataacc ataaattgga ggattntggtt acaattatgc cagtttatac 480
 a 481

<210> 954
 <211> 436
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 954

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 aactcaaan atgcagtcac atatgtacaa attntttcca caagctaaat tccatataaa 120
 cacacgcaaa tgccattgag gcatgtcacc gaacacttga tgggtgcatg tttagacatg 180
 aaaaaataa ggaacgngg gaatgtgaca tgcccattca tctcagagtt cacaataggc 240
 ttgcgccat ccatacaac cccccaattc aaacaaacaa gcatgaatcc aaacattcat 300
 ttctcatga aatntgaaaa tacaagcaaa caaagcacta aaatacagca atggcaagcc 360
 aaagatcana ggagaatgac acttaattgt anggagtga acaaaatgca taaaggagaa 420
 caaaaactca acaatg 436

<210> 955
 <211> 384
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 955

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[illegible]

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 ttaatgacgc acggacactt atgcctgacc taactacgat tcacgcaaca tgtcagcgca 420
 gatcaatacg gataangact gatatggact ggatcatacc atntgatcgc cgacngacac 480
 gtaatacacg tggagcctaa ggggctacat gcatgctgcc tacga 525

<210> 958
 <211> 523
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 958

tctaattaac ctacatatga gagaagatga ttattaaaca ctcataatga agataactaag 60
 tattttattac ctataacttaa tagacaatac ttataagctt acaaaatagc catatattga 120
 gagagtgtga tacaatttat acaagtgtta tacgtataag ttagatgttc tcaccgacta 180
 acatcctatc aaggtcatca aaattagacc agtttccatt cttgaatgac cctaacaaag 240
 catgcatgta cgtgatcaag gtaaaggcat actagaatga atagctgata gcacagagaa 300
 cacaccaaac atcattaaat agatagaatg atatttacat caagtaccta caaggaagat 360
 ccaacagagg attntagctt tccatatcca ggaagccttc tttacaacan agagaagaat 420
 aagatgacag agtgctgcta tacaagcggg gaggatgtct tcttcacctg taggatctca 480
 caaccactca agaactcatc tcagactcat agaaacggct tcg 523

<210> 959
 <211> 304
 <212> DNA
 <213> Glycine max

<400> 959

atcggtgaga gtgtaacctt aaactgtgag tgaacgacta gctgtgagta ataatctttg 60
 catgaatctc tgaatttttag aatgaaatgt ataactgaga acatgatgaa ggccatgatt 120
 gtacatatatc aagctctttt gaccaaacia cttaccttga atgataattg catcctttgc 180
 tccctttttg agctgaatga tggtgtaaaa aatttgaacc ctaaactaaa ataattatgt 240
 cttgatacct tggttagatt ttaggagagc atatggttca aggcaaactc actctaaatt 300
 tggg 304

<210> 960
 <211> 344
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 960

tctggtggga catcttgact tgctntccaa tctgacattc accacanatt ctgccttctt 60
 ctattttcag aatgagaatg cctctaacaa cacctttgtc aatgattttc ttcatgcctc 120
 ttaagtgcag atgtccaaat ctttgatgcc atattctgac ttcattcttct ttggaggata 180
 gacatgtgga ggagtaactg gtttcttgag gtgtccatag gtagcagttg tcctttgatc 240
 tgctgccctt cattagaact tcactcttct catttgtcac caagcattct gactttgtga 300
 agtttacatt gaatccttca tcacacagct gactgatgct gatc 344

<210> 961
 <211> 385
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 961

gtccatgagg aatctccttg ggaaagacat ctttaaattc ctgcaataag ggttgaacac 60
 taggagaaac ataaatagtt tactgattag aattatcact ctctctctct tgtgtatcac 120
 tccatctctc aagtgtatca ctcttccttt ttctattcct ctgtgatgcc tcactattgg 180
 ccctctcttg gtctctcttt tctctccttc tgattcggac atcacacact tctctgaggg 240
 ataaagtttt atgaataatt ttctggatcat ggtgctggag agaaatcttg tttgagaacc 300
 catcatgcac tgctttggag tcctcctcaa tgatggccct cactagtgcc atctccatct 360
 ncttatcact aacaatcaaa ctttc 385

<210> 962
 <211> 247
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 962

ggtagaatgg ccagacatga tacatgtcag ggattggttt ggttcatagg taatatggat 60
 gtccacatt atttccatga cacatatgca aaaatgatga cttggatact ntatgccaaa 120
 ctggatcatgc atgcacctat gtggacactc atgtgtcaaa tttttatggt catgtgatgc 180
 tagggctcag gattcattct ctctatttta atcaacccaa tgtttccaaa atatgttctt 240
 ttatcca 247

<210> 963
 <211> 357
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 963

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 tggcataatc aaactaaaat aaataataac caagagtcaa taatcctttc acgtgtcatg 120
 aggacgataa acctgagcat cattaacatg gtcctcagtt ccatctgtta tggtagtgga 180
 tggactanga gctagcatga tagactgaat atgcaagtgt ttagtatat gctgcatttg 240
 ctcatnttgt tgacaaattt gctcatnttt tcttacatct gttgtcgcac ctcttgactn 300
 tgggtggcat gagtctccat tagttgcaac atcttttctt caagtgtctt ctcttta 357

<210> 964
 <211> 406
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 964

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 tgcagaaagc gctaagcacc gctgctgcac taagcccaa atgcttattg gaagttacaa 120
 cttcaagttg ggcttagcgc gngctagc taagcaccag tgttttaaac tcaaagtca 180
 tgttggcagc ctaagcgcac catacgaatt tcagttnttt aaaagtagag gcagaggcac 240
 ttgggttgct accttggcac ccaaacctct gactctctc atctctgagc atatttctat 300
 ttctgctttg tgcttattga tctctacat ctntcttcac actctgcatt acacaatcca 360
 agtaagtaac ttgatttctt ttcactttta ttttcatggt tcacag 406

<210> 965
 <211> 330
 <212> DNA
 <213> Glycine max

<400> 965

aaaaaaaatt aaaaattaaa accagggttaa gggagcttac ttggtggtga ccctttttca 60
 catttccttt ttcccaaatt tggaattact ctttttatgc cgaagccggt taccggaaag 120
 tggctcggat cggccaagta ataattaaaa cggaatgata cgagtgtcaa cacagggaac 180
 ttattccttt ggcaaagctt tgttcaacaa tcatgcattt tggtagacaga aaataataat 240
 tgtgaattga agtaaaagta tgatatatcc taattgaaaa gcagtaaacg tgagcaaata 300
 agtgtgaaaa cagtgatcta aaagcattgg 330

<210> 966
 <211> 421
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 966

tgattntngt tctgattgca tgatgctagg atagttgata gttaanatag tgtaggaat 60
 tatattttca tacaatgtat gttgttctgg ttaggaattg atggcctaata ttagaagca 120
 agcttcatga tgatgaacct agcaattttg acgatgcaa aagaccaagt gattgattca 180
 agacttcaag atcaagcaca aagaatctaa tccaagattc aagattcaag agaagaaatc 240
 aagaagcaat aagtcaagac ttcatatagg ataagtatta aaagaatttt tcaaaaacaa 300
 aatagcacag ttttgggtata caaaagaatt ntctcaaatt ntttaagtta ccagagtgat 360
 tactctctgg taatcgatta cctgttatca gtaatcgatt accagttgtc ataccctaata 420
 t 421

<210> 967
 <211> 488
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 967

caactttccc aatctactcg aagcacaaga ttgtcaaagt tgggtggaaat gaaaggagcc 240
 tttcatccac agctagtcaa gggtttctac agctatgcac ttgttgaccc ttgaagtaac 300
 ctctcttcta aataaatgga gtaaaatagt cttgatctat tgatggaaga agtactggct 360
 ggatatgggg gagtcacaat tctataaacg atgatgggac aacagatgag actatganga 420
 tgttctgacc tttagaactg aaaatcctaa agtgtggctg ttgagagcaa atgtgttatt 480
 ctt 483

<210> 970
 <211> 376
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 970

ntgtcctcag atcactcttg ttggactcag cccaatcgag atactcctct taggtttaga 60
 ctaacttaaa ctgagtttca tctgcagatc cctcttgtag caatgccaaa tagtactcta 120
 acaggagatt ctttggtgga agtagtggat tataccatca agacaagaga gcagataaca 180
 aagttaatta cttcaccaca acttgcttcg cgcccaggaa agaatgaaac attatgctga 240
 cctcaaacga gtggacaagg agttcaaatg tggagaccta gtctatttaa taattcactc 300
 atacaagcag cttactttgg canactatgc tttccacaca tagcagccac tagcgggtcta 360
 tgtaaagaaa catcat 376

<210> 971
 <211> 338
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 971

ngttatatac ggaagacaag aaaatgggat atgaacgaaa tgatgatgaa agcttagaat 60
 ctagacatga attgaaagtc tcagattcga aaacttaccg gttgaataat gaagaacgaa 120
 tgaagaatga atgaagaacg acggaaaacc atcatggatt tgctcacgat aacgtctcgg 180
 aagcattaca gaagcacctc ggcttggatt ttcttcacgg aaacaatttt ttttcaccag 240
 aacagctgaa atgcatagcc aggggatccg ggatccttgg aacaaccccc tttttctctc 300

338

<400> 972

<400> 973

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<223>      unsure at all n locations
<400>      974
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400

tatcttcatt tcaagaagaa ttaagtgatg aaaatataag gttggcttga tgataaaggg 180
tgaaggaaga gagggatagg aatcttccgg ttaacaaaaa ctaataaatt aacaactaac 240
atctaccgat aaaaaaaaaa aagatagaat taagtgatta gttttatctc cttttcacat 300
tataggttct tcaatgtttc ttcacaattt ctaaacatt gtctgtagac tgctccaagc 360
ccagttcett ctcaaacagc cccagagaaa tgacattgat gtgtctaata taagttctct 420
cttagtctta aaatttcaaa tgtagtgttt actatac 457

<210> 975
<211> 577
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 975

attgacccat gtgaangcga tctattangc gacactacag aagactcagc tcagtcacac 60
tagtcggcag acttatctga atcttttgaa gatgagagaa cagccaatt aanaagagaa 120
agggagaaac ttcaagaaca tgggtcttcac tcttcacgtc caagagaaga atccccacaa 180
atatcacaaa caactcttcc acggatttca ccaatccaag cactcacgtt caaagcaaatt 240
aacaagagca acagaatcaa acccatcatt gtcttttgtt tttccatggg tgagattgaa 300
aaaaaaacaa caaanaatgg tatctttcac aagcactatg tctttggatt ctatgagtag 360
ttttcttata accataatgc tcctaaagga tgtttatggc tcatagtgc tggaattnttt 420
caccgacatg atatcttata ttatgtgtaa gactatctga actntatgtt aagatcactt 480
ggattctcaa ttttaatgga tggttntagc taataaaaga atcttatatt gtacgcctac 540
aaataaaaaga acaaatcgtn taactttaca ttctttt 577

<210> 976
<211> 460
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 976

agctnttagc attntaccat attatataat ggctagcttg ttatatattc tcctaactac 60
atggcatgcc aagatatcac taatatgttt aatgtccacg gttgtttctg ttaagtttgg 120

gctacgtgag ttgactctga taccattatc attatcatga tgtactgttc atggacccaa 180
ctatatatta tattattaat tactaaaata agagttagtt ttaaaacact acaacttttc 240
tactataaat gtattggtga aatcattcat tgaaatgata atgctactta catatatata 300
ttatttaaaa atgtatatta taactntacc acttctctaa aaaaacaaaa aanaaaactt 360
ttccatttta cataaatatg taatatgggtg ttaattgata tgtatcaagc atttattttc 420
actaacccaa gtctgcagga attctatgta ctttgcatt 460

<210> 977
<211> 307
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 977

tgcataaatg agaggaaact tgagtaagtt ntatttataa atatatatta cagatgatca 60
tcttttaatt tcgaaaaatt atattcactt ttattctgca ataaatgttt agttattttc 120
tggtgaatta gaatttaatg ctgatttttc aattaatcat ttctaaacta atatcatggc 180
ttgtggacat tggaatctat tacattagtt tccccacgct aaaaaaaaaa tagtgatggt 240
caacaattgc ggcgttattc catgtgcatg gagatcaata caccgacaaa atagtgaatg 300
caccat 307

<210> 978
<211> 260
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 978

ctttgatagc cctttntgag ccttgtttcc ctttccttgt tttgaagctc actacaagcc 60
ttaagtgaat aaccatgata tcaccatata ctttaaggaat tttggagctt tggaattggt 120
ttgggaataa gtgtgggggg tttttgtttc attggacaac ttgttttggt ggctatgctt 180
catgatgtat tctgggccat acttgatgta cattgtatat tggttaaatg ttggacatgc 240
tgaatgaaat gttgttttctc 260

<210> 979
 <211> 401
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 979

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 gcttgcttaa ggtccaaaag gaacttgact cagcgtttgt gcgagacaga gatcaacatg 120
 ttggctatca tcaccaagta tcaataagaa ctaagtctag ccacagccca cgagcatagg 180
 gtggcaaacg agtatgcccc agtgtacgcg gaanaggagg ctagaggaag ggtgatcgac 240
 tcgttacacc aagaggcaac catgtggatg gaccaatttg ctcttacctt anacgggagt 300
 caagaacttc cccgattgct agccaaggcc aaagcaatgg tggacaccta ctccgcccnc 360
 gagggagatc acagacttct cgactattgt cagcatatga t 401

<210> 980
 <211> 363
 <212> DNA
 <213> Glycine max

<400> 980

gcagctaact accatgcact acggatatac tctaaggaac gcaaaaagta acaacaaaga 60
 ccacattaag aactacatat gcagcgacct caacgcaaaa gtattacctt cactcctcgc 120
 aaccaaaggg aacaacacac caaaggaaaa atggaaaaca aagggaactt atccaaaaac 180
 aaagcatgaa agtcagcaaa caacaaaaaa ggacgcgaaa agatagaaca agaaggaaaa 240
 atttagtaaa gaatcataga acgtgaagta aacaacaaaa caataacaaa ccgggtagaa 300
 aaacagctac agccgcgcca aacacagaca tgcacaacag acgtaaaaaa cttacacctt 360
 gaa 363

<210> 981
 <211> 366
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 981

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attntccact atggagatgc agcgaaagac aaaggagaag aggtgagagg aggcgccatc 120
 cactgtggaa taagccatga aagaaagagt ttcaccatca agatgagcct tggataagaa 180
 gcttggaagg atgcttcaat ggaggaaaag acagagggag agaaagagat aggggggagca 240
 agaaatcgaa ggaataaaaag agggagaata gtggaacttt gaagtatatc tcacaagact 300
 ctcatcctac anagttacaa caagtgttac gcatgcttct atntatagac taagtagctt 360
 ccttga 366

<210> 982
 <211> 514
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 982

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 gtgcggtcta ggacacaatg tcaattcata cgatatgcga ggatgactcc ccgagcaagt 120
 tggatttggg atgaccatgc cctcctggtt tctgactang aaattggcga gtggaggagc 180
 gccacacat ttacgcgaca agcataatgt aaccctttgt ggctnttaaa ctctacggng 240
 gggcctangc tntagagatt ccttttggtt tggcattatg tcttttggtc ttgaatttat 300
 aaatataaag atctttcttc atctgttctt gcacctctac ccattctcat tcatttgcac 360
 gtntatttct ntacgcttaa nacactagat ccaacaacga gtccctcnaa ggtactaata 420
 cctgngaccc tgnatcgat tcatgcaaga agcgggcaca cagagagtga gaggacgatg 480
 atgtgtactt tcccacagtn gagaaatagt actn 514

<210> 983
 <211> 369
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 983

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 ccatctcaat atttacgaga tcatattcat acaccattgg ggcatttcac caagcacttg 120
 gtgagcgcat gtttggacat gaattgcaag agaatgggag caatgtggca tgccccattg 180

cttcaaaata caacctatgc ctaagacctt ttcattcaga ttctcaattc aagataacaa 240
 ggcctaagc taaccataac tgcctcacia atataatgca tgttctcaca atttagggca 300
 ccaaaagatg aagaaaacac atcantggga agcatatata tcaaagatcg aatacttact 360
 tgttgaggt 369

<210> 984
 <211> 419
 <212> DNA
 <213> Glycine max

<400> 984

agcttcgctg atttagtttt caccgacgaa atgatcgaag tgggtctaaa aagaggcaaa 60
 tctgatcatc atgctttgat aaatgcaaaa aaaattgggg caagtgaaga gggtgagaat 120
 gaaggagaaa cccatgttgt gactgccatt cctatacagc caagtttccc accaacccaa 180
 caatgtcatt actcagccaa taacaaactt tctctttacc caccaccagc ttatccacaa 240
 aggccatccc taaatcaacc acaaagcctg tctatcacac ttctaataac gaacaccacc 300
 tttagcacga accaaaacac caaccaaaaa ggaattttgc agcaaaaagc ctgtaggatt 360
 caccctaaat tccggtgtca tatgctaaac ttactctcaa atctactcaa taattcaat 419

<210> 985
 <211> 423
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 985

nggatggcaa tagaattaac caattattag tttggccata ggaagttcga ttgccttatt 60
 ntatttcgat tntgatttca ttatgtttct ttaattntgt ttcttacgta aacaaagcaa 120
 ctgccaaatg catttccttg gaacaaattt atttgaatat gtctttgggt ttgttaaattc 180
 aattnttaac tnttagtaga cttacacaaa tatgttatgt tatccaatgg gcatatgtaa 240
 cgggtctaaa gattagacaa cgtatattct tcattaagaa aaaggaaaag gcgattataa 300
 ttntgactta agaagttggt ntgattctgt ttactgattc anaagttggc tgattgttn 360
 ttagttntct actggatttt attatccttg aggacttggt gtgtcctcac atagtgtatt 420

ttc

423

<210> 986
<211> 311
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 986

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aaagttatga ccattagaat ttctcgagag ctccggttgt tcaatttcaa gagtctagat 120
gagttatgta cgcgaatcga acatctgtgt gaaaagttat gaccattcaa atatcttgag 180
tgcttccggt gtgcaatttc gagcatcttg atatattatg tcccacattt ggacattcgt 240
gtgaaaaggt atgaccattc gaatttctcg agagcttcca ttgtttaatt tcgagagtct 300
agatgagtta t 311

<210> 987
<211> 575
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 987

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cngcgtttcg ttcatgtgtc tccaccttcc gagttggagc tatgcgtagg gattgcttag 120
tgcaattctc cattctcaac cttntcggga gcccattgaa ttgcgttttc gttcatgtgt 180
cctccaccct cgagttcggga gctatgcgta gtgattgctt agtgcaattc tccattctca 240
aacttttttg gagcccatg aattatgttt tcgttcatgt gtcctccacc ttcgagtttg 300
gagctatgcg tagtgattga ttagncaat tctccattct caacctttta cggagcccat 360
gaattgcgtt ttcggtcatt gtgtcctcac ctccgagttt ggagccatgc gtagagattg 420
cttagtgcaa ttcttcattc tcaacccttt ttcggagccc atgaattgcg ttnntcgtca 480
tgcgttctcc acctctcgag ttggagctat gcgtagtgat tgcttagtgc aattctccat 540
tctcacacct ttccagagcc catggattat gtttg 575

<210> 988

<211> 415
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 988

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 gaatctctgg atcttgattc atccactgat ttacctttct catttaagtc aaggtagggt 120
 gatgtagcca taagaatgga cgcttctttg cattnttcca taccaaattt ttttattagt 180
 tctatgcagt atttattttg accgaggaag gttccatggt tcattttctt gacttggagt 240
 cctagaaaga aatttaattc tcccatcata gatgtctcaa attctttttt gcatacaaca 300
 tgaaaatccc ttgcataagg ttccattagt atagccaaat ataatatcat caacatatat 360
 ntgaacaatt aacaaatcat tgtttacttt cataatntaa caaagtttgt caact 415

<210> 989
 <211> 260
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 989

cgatgataaa gactcccca gctatntatc ttctctctca gagaggctnt gtctcactct 60
 aagaagtgga tcaactcttat cttggatgga taggaatgaa agctcctaca cttatttata 120
 ctactccatc tncacaataa atgggtggaga ttacttgtct cataatgtga agattaattc 180
 tctataatgc ttcacacatt ctaagagttt ctacactctt ccatattctt tcataagggt 240
 ccagaaagtt ttacacatct 260

<210> 990
 <211> 181
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 990

tactttgtca ggtacatggt cacttgcaat ttcatctctt agctccttgt cccttgtcaa 60
 gatcttccaa gatactatta taacttcttt aggaatctct tctgaagcct atccttcaag 120
 ggtagcagcc ttcttttttg cttcttcttc tgctntcttc ttcatggtct tttatgctct 180

t

181

<210> 991
 <211> 585
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 991

agggtacaca gcacaaaagg acngctgcgt aangctacct atcattactc aagctcgaat 60
 gaggaagtgt agaacggtga aacttcctgc tnttattctt tgaccacaga gtggtacctg 120
 gagatatgtc gcgggtatag tcagtcagtg agaacctgtg atgtacctaa gcaggcgagc 180
 tcctggcagt caacagataa aaggaacaaa gatcaciaag caaggaggct tgtgtggttg 240
 ctggccagtt gtgaaacttg attgatatat gggatgtggc ctctggtaat cgattaccaa 300
 ggggtgggtaa tcgattacaa ggcttanaaa gtgaagacag gaagctaaga tggcctctgg 360
 taatcgatta ccaaggggtg taatcgatta tcangcttga aaatgggatt aggaagctaa 420
 gagggcttct ggtaatcgat taccaagggg tgtaatcgat taccangctt anaaatgaan 480
 gcagcatgtg gtggaggcct ctggtaatcg attaccaggc tgtgtaatcg attacacagg 540
 ggaacatgcc actggtaatc gttaccaggt atgtgtaatc gatan 585

<210> 992
 <211> 320
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 992

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 attccttgct acgttaataa gcataacctt gtatgtctcc atcttgatgt agccttncat 120
 tgattcctca aacttgacga ctccccata tttagtcttg caacttgctt tcaaactgca 180
 acatcaatga tcattctctac atcattaaca gtagagtata aattgccttc catatcaaca 240
 cgggcacatg tgtagaaaac ctttaccac tcaaggtaga agattnttat cttctacacc 300
 aatcttaaca aaccatatgc 320

<210> 993
 <211> 126
 <212> DNA
 <213> Glycine max

<400> 993

tatatcatcc agttccagtc atggtatata gtataaaaat ttaaacaatca acacaacatg 60
 caatgaagcc tagcttccaa agacaacaag gttagggttc aacaagtgga aagaccccc 120
 cccct 126

<210> 994
 <211> 248
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 994

attcccttag caatccccca aattaagaac ttatcataac ttgaaaccct tatactctct 60
 tagaaccta aaacaagatc aaggttatca aaattaggct caggggttta ttcaaacaaa 120
 tcattattac ttttggtca ataggggtgc aagggataaa ttcatcacag gttggctntt 180
 tggctgagtg gctaanataa aaagaaacna tggcttgatc atatccacct tatgcaaata 240
 atcaaata 248

<210> 995
 <211> 305
 <212> DNA
 <213> Glycine max

<400> 995

tatagatact cagcctcatt ggagcttgag gcctaggatc ttcttatcaa tggattctct 60
 tgcttcttgg aagatgaatg gaagcgaat ggagaaagg agagagagag gagacgccac 120
 ttcaaagaga agatgagtct agaagaagct caccaccata ggaggccatg gataagagct 180
 tggaggaaga aggagatgaa tgaagggaga gggagagaag agcacgaaca tttgtgctct 240
 acatgagctt tgagatctga agtttaatat tcaaatgatc aaagttgaaa aaaatgcaca 300
 cacat 305

<210> 996

<211> 362
 <212> DNA
 <213> Glycine max

<400> 996

gcttggtggc cgcgattgac aaaggggtgca tatatacgac gttagtctct gcatgctatc 60
 atgcgttgac tgtagcgat agcaaaagaa tgtttataact aataaccact tgggtatttc 120
 tgccggcccc ctaacttcac gacttagtac cgacagagtt tgtaagcgtg gaagacgacg 180
 taaatctccg catgtgaacg agcttggttg ccgcgattga caaaggggtgc agaagacgac 240
 atttgttttt tcatggatc atgcattgag tcttagagat agcaaaagaa tgtttatagg 300
 gataaccact tgggtatttc cgccgacccc caacatcacg agtttgtatt ggagagggtt 360
 tt 362

<210> 997
 <211> 416
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 997

nttaggacct tatgtttagt tnnttggtt ctaaattctt tgacttgtaa acaaaaagtt 60
 tctttgtaat atttcaatgc ttaagtgaag tgttcagatt atgatgttta caattacttc 120
 aacaagtcct tcgataataa attgttcttt ctttttgcac aagcataatc atgcattcatt 180
 ctgcattcat agtttccgca tcaagtctca cactgtgttc accacttcaa aaggataatc 240
 agccgcccgc cgaagaaagt ggcccgcga attctccgca naccaactcg cattttcaga 300
 aatggatcta attgagcaag aaaatcagag tctcaaggag gaggttgcca cntacgaga 360
 aggaatggat aggttgacga ccatgatgaa tgcactcctg tccgcccgaga attctc 416

<210> 998
 <211> 435
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 998

agcttgaggc acttgccctt taacctagtg tctccaaagt ggccttacc aagtatcttg 60

ttttccattg atgtgccatc attttcttct attttctaaa ccctttttgc accattttta 120
 ttattgattg gtcttaattg tcaattaatt aggcagtttt attatttggg cccattcagc 180
 caatttgatg tttttaatct aatttcagga attaataag aattgggctt gaatctagca 240
 ttgggcttga atctagaatt gggcttggac ttgaagaggg caaactaatt tattctataa 300
 aattagatct tatcttatct agatattatt tagatttgat ctcatctaga tatcatttca 360
 attagatctt atcttatctt atcttatcta gatntgattt gattntactt atgggcttgg 420
 atttaaaaca tattt 435

<210> 999
 <211> 243
 <212> DNA
 <213> Glycine max
 <400> 999

tgacactact tatcttacc tacttctacc accaaaatta agtataacct atagaatttt 60
 actctgaat taattaatta aacgaatgtg tatagaactt tctattttct tttcataagt 120
 aaacattcct cgcttagacg ctctgctatc ttcatagcgc tccaatcata ttaatagtta 180
 ttccacccca ttctgtgata tacaacctga aaagctctga atatgcttga tacggaatta 240
 gtt 243

<210> 1000
 <211> 257
 <212> DNA
 <213> Glycine max
 <400> 1000

taatttatct caccctcatt tgtcacaaga tagtgacatg gagttgatgg tcaatatttg 60
 tcacaacaaa gtatgtttat ctcaacctaa ttgttgcag cactccattt ctatatatta 120
 caattattca tgttcggcat tagcatgtac gtccttgcga ctattgttcc acccatagca 180
 aggaataagc taaccataac atgagcccaa caaaggaaga atgctgatat agatgatgca 240
 gtataaagaa aactgaa 257

<210> 1001
 <211> 522
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1001

ntanatatnt cagatgaagt cattcaatta ttgttacttc atatgtagca gctgggagga 60
gatttatatg tttgtcaa atgtgttgaca atgtagaaac attgttaggc atgacaaccg 120
gatcgggttc tgcttatcct atctcagtc ccgattcctc atttcttct ctaccctgt 180
ccccgaaatt caatgggagt gcatatztat gtccatccca gtctccagtg gggttgagtt 240
tttcccgctc cgtcctgccc cggacatatt tataaaattn tattaataaa tctaattntt 300
cataaatga agaataataga ttttaataaa aatcacaata ttgtacatga caacatanaa 360
tccaattcaa cattagcata naattcaata taataatttc atggttta atgtgttatata 420
tatatatata tatataatga tatntttgta caataattat tagcgtggag gaattggagg 480
cgggtattaa taatctcatc cctgaccccg aacctgattt tg 522

<210> 1002

<211> 267

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1002

ctcgtgcan aattcatttc ttgggtgttg ctcttggtt gtgctaaagg tgggtgttcgt 60
cattggaagt gcggtacaca gactttgttg ctgacttaag gattgccttt gtggataact 120
gggcgggtgg taaggagaac gtttgttatt ggctgaatca tgacatcggt cgggttggtgg 180
gaaacttggc tgtctatgaa tggcagacac agcatgggct tcatcttcat cctcaccctc 240
ttcatttgcc ccagttttct cattcgt 267

<210> 1003

<211> 360

<212> DNA

<213> Glycine max

<400> 1003

tattccgagg acagttcatt atcatgcaca gcctgcaaga gttggctcac aatacgccaa 60
tcatatctat ggagcatttt ctgagccag tagcctcgcc tgaagctcaa cttccattgg 120

tgaaaccaaa cgaggttgct ccgcctgagc tcacacctga gtaggtcaat tcagagccag 180
 ctaaccaca atctccagtg gcgaatccac cttcttcgct tgagcttgaa gcaagtcgcc 240
 catctctcc tctgaatgtc atttctgacg catcattaga tgaagcattt gtccttctg 300
 atttaccagc tgcagataca gctgaccacc ttgtttcccc aatcggagga catgctgac 360

<210> 1004
 <211> 237
 <212> DNA
 <213> Glycine max

<400> 1004

agcttgcttg agaagcttct atgtaagctg gatctttgag cttcaataaa ttccttcaat 60
 tgtgattttc agccatggag ttgcagtgga agataaagga aaagagatga gaggagacgc 120
 catccactag agaataagac atggaaagag aagcttcacc accaagagag tgtcttggat 180
 aagaagctta gagaggaagc ttcaatagag gaagagaatg agagagaggg agggggg 237

<210> 1005
 <211> 243
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1005

cttganatcg aanactaact ggttgaagat ngacgaacaa tgaatatcga tataagaatg 60
 gtgaagaaca ctagtataat tcatcacgaa aacgtcacga aagcatctcg gcttggatta 120
 ttttcttctt tcttcttctc ctactaatt gtaagtgaat tttgagtgcc aaaggtggtg 180
 aacccttttt cctcagcccc ccatgccatt ttattgaaaa aattgagggg gggggggggg 240
 ctc 243

<210> 1006
 <211> 315
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1006

gcttctcca caagcaacag ccttctggag gaatcttctg gaaggcccaa gtgggcctgg 60

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tattatgggtc taagctgggt acacgaacat ntattcccat catatataat aaataaaaaac	60
aaaaattgag ctccaccaa aacaatagtt tgaaaatcaa cttcactaat gttggagaag	120
tcagagaaaag ttgctagaca aaaaatatct ttgtcatata gctggaatga tcatccgtga	180
gagaaaattc atggcaagca tagcaagggt cattcttctt gaaaccacaa ttggaagctt	240
caaactaata atcttggaga ccatattatt taagcttcaa attaatgcat cttgaactta	300
gtcacgacta gtacactaga gacacgataa agtcaccaga naacacgcaa aagtacttaa	360
taaanactta taagttaatg gttagattaa taacatttag tttaacatgg ttttaggttg	420
tcaaattggt ccgtcaaaaa agtttc	446

cgcatgatac	atctgacccg	cgatctctga	gtcaacttgc	agctgcgcag	cctgggtggcc	60
catgaaggat	ggcttgcggt	tatattgcat	gaaaaagccc	ttcgattata	tggatatatg	120
tgaatgggta	gcataaaatg	gcttgcgaaa	tgggtgaataa	aatggccttg	caaatatgaa	180
tatatattgc	ctggaaatgg	cttggattat	atgaatatat	attgtatgaa	gtggcttacc	240
aaggggtgga	tggatagccg	aaaagtgggt	ttcaaaatat	gtggatttgt	gaagaggagg	300
caaaagaagc	cttccaaaaa	aatgtgtgat	atatatagga	tgtaacgtga	aagggttgca	360

aaaaatatga catggatgtg tgctgaaagt gctttcacaa attttatgtg tgcaatgata 420
 tgtgtataaa atacatggcc caaatgtgat ttataagtgc tgtgacactc gccccatgag 480
 tgtgtttgct cttgttg 497

<210> 1009
 <211> 441
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1009

tattaagaag cttcctccag aagcttcctc atggcttctn tgagaagctn tctcaacaag 60
 cttctttgag aagctagatc cttatctatc cacaccctc tattaactaa attaacttcc 120
 ttaaaaaataa ttacggatga aaataatgca acaataatc aaacatcaaa cataattact 180
 aataatatat atatcagggg gttacacatg gtatacttga gaccgtatag taagcataaa 240
 attgagtata ccaagaacaa tgccttttta ttgactacaa ccaaagctat aagggtcgcc 300
 aatgataggc actaagttgt aagatcaata tttctatata tgttgaattt caagagttgt 360
 agttcctttc taaactanga acaaaanana aaggataaaa aacatgccac ccctctaaaa 420
 tatcacacaa ctntntttaa a 441

<210> 1010
 <211> 444
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1010

agcttctagc caaatattag tctctcaaag agtcgtcttt tcatggagga ttatctgttt 60
 ctagagtctc atcaattcga ggtgtccttg gttttatgga gactcatttc tatttaacta 120
 tcttgagtg ccaatctttc ctggaaaatc gagcaaaata tatctccaaa gtgtgttttg 180
 atgagaaaat tttaaattct gagaaatttt aaattctaag aatttcaaatt acttcaattg 240
 aaattctttt atttttaaaa ttgtgttttg ataaaaaaa ataaaaattg tgaggggtgaa 300
 agaaaatgaa tgcaaaggga agagaagata tgattggtgt gtttttaaaag agaagaatat 360
 tgacacggca tggagagtca caganaact gggacacgac gacatacacc accatacccg 420

accacaacat tcagtcaatg acac

444

<210> 1011
<211> 470
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1011

tactcaagct tcaacaacga gggtcagtga atgattacct cactgagtct gaacgttttag 60
cgaatcggat tgttggtctg tctccacagg tgctcctaag ctgttttatt tcgggccttt 120
accagacct ccgtcgtgaa gtccgggctc tgcaacctat gtccatatca caagtcgtgg 180
cgéttgcaa gttgtaggaa gaaaagattc aggaccgcca tcgccatttc cgcacatcct 240
ataccccttc tgggtccgca ctgtcaccgc caccaccac cgcggttcct tccatcggtc 300
tcaccccggt acgccttca gttaagcgcc tttcagcaga agaacttggg gtctgtcgtg 360
acaaggggtt atgttatcat tgtgacgaga agtggattct cgacaécggt gccgtcctcg 420
cctccactta cttattgcan acgatgatga tgatgactgc acaaatccat 470

<210> 1012
<211> 417
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1012

ttggcttatt ccctagtgga tgggtgcctcc cctctctct tctcctttgc cttgcgctgc 60
atctccatgg tgtaaaatca ccattgaagg acctcattga agtcaaaga tccagcctcc 120
atagaagctc cacaaccaag cttccatcag gacgaagttt ggattgattc aatctaacta 180
gggattgagg tttagtaatt taagctatag catagaacac aaaagcatga tngattagag 240
aaacatcttt atatacatca gttgggttgg tagaaagact caacatcttt acctactggc 300
tgcaatctta cttactttgc attttactgg ttttagccta gatntagtnt aattctattc 360
taaactatcc attatcaatg gttctctcac aatgacttat tctgaattaa ccctatc 417

<210> 1013
<211> 391

<212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1013

tctcaaggga agtttcttaa gaaagcttct caaggaagct acctagtcta taaatagaag 60
 catgtgtaac acttggtgta actntgatga atgagagtct tgtgagacac aactcanagt 120
 tcaacttctc tccctttgtc ttccttcaat ttcgtgctcc cccctctcta tttctctccc 180
 tctttctttt cctccattga agcatcctct ccaagcttct tatccaaggc tcattcttgg 240
 ggtgaagctc cttcttccat ggcttatctc ctagtggatg gcgccgtctc ttacctcttc 300
 tcctttgtct tccgcttcat ctccatgggg gaaaatcacc attaaaggac ctcatgaag 360
 ctcatagatc cagcctccat agaagctcca c 391

<210> 1014
 <211> 332
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1014

ngaccaatch cgaccaacc caggcatagt cggtcagtga gaactctgtg atgtacctaa 60
 acaggcgagc tcttggcagt caacagatca aaggaacaaa gaccacaaag caaggaggct 120
 tgtgggtggc gccagctgt gaaacttgat tgatatgtga gatatggctc ctggtaatcg 180
 attaccaagg gtgggttatt gattacaagg ttaataatg aaggaggcta acatggcttc 240
 tggtaatcga ttaccacggg gtgtaatcga ttaccaggct cgaaaacgag gtcatagaag 300
 catgagggct tctggtaatc gattaccagg ct 332

<210> 1015
 <211> 348
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1015

gcttctatgg agctggatct ttgagcttaa taagggcctt caatgggtgat tntcagccat 60
 ggagttgcag cggaagataa aggagaaaag gtgagaggag gcgccatcca ctagagaata 120

agccatggaa ggagaagctt caccaccaag agagtgcctt agataagaag cttagagagg 180
aagcttcaat ggaggaagag aatgagagag ggagagagag agaatggtgt ggaaattgaa 240
ggagaatagg gagataagtt gaactttaaa gtgtgtctca caagtttctc attcatcaaa 300
agtatgacaa gtgttacaca tgtttctatt tatggcctag cacatggg 348

<210> 1016
<211> 375
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1016

cgaagaagtt ntttctttta catgccaac tctctcgagt gacattngca ttgattggtg 60
tattttatgt tgcattcttag tctctatcat atcctatgtg catcatgcat catcatgtgt 120
gagtaaggag aaaatttcta atgttataaa atttcttcag aaggcaaaac tttttggttt 180
aatccattac aaccttacag taatcaatta caciaagttg ttttaagcttg catagctatg 240
tcttgatcg atttaattaa ttacagcctt atcctaateg attacaccaa ttgttttaag 300
acaatggttt atttatntaa tagtctatgc ttttaatcaat taccatgtga tataaatcaa 360
tacttctctt tctat 375

<210> 1017
<211> 400
<212> DNA
<213> Glycine max

<400> 1017

agctataacc tcacgtccc tcacagtctt tattattggg agccaatcca atccttgtgt 60
tcggactctc agccacttat gatagccgc gatgatccca ttactgcttc ccctaagctc 120
tctgtccttt cttcacgccg catcccatgc cttgcgaact cttggagta cctcgcgtt 180
gtggtcacta aaaccccggtg cgatgaaagg cgtgatgctt tcgtctaata ggcctcctct 240
catggggtag ccaagctgtc ttatggtgag aacgggatta taattaatac aacccttgt 300
tcccatcaag ggaacatttg gacatccttc gcatgaagat agaatcctga ttcttccttc 360
cttctagcga gggaaccaat taacagacgc ccccccattgc 400

<210> 1018
 <211> 267
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1018

tattacaagc ctttaagtga aaaccatgat ttcacnctac ccttaaggaa ttntggagct 60
 cttgaaaatg tttgggaata agtgggagag ggggtatgtt tcattgggtg atattgtttt 120
 cgtggccatg cttgatgatg attttggcca tgcttgatgt atatacatat aatgcctata 180
 tgggtgcttta tattttaaat gctttgcaat gctactggtc acgttcaata aaaaattaaa 240
 tagaagaaga atgatgttga ataaatg 267

<210> 1019
 <211> 389
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1019

gcttgcgagt ctcctttggc atttccttga catatgttct ccttatgtct aaggtctcaa 60
 ggttatgaag tttgccaata aatttcagaa gctgcgttgg catcactgaa tatcgtaagt 120
 ttaaatactt gaagtgtgct agatttcccc aattttcagg aacagaactc aatggactat 180
 cttgaaaatc aagtaccttc aatagcctgt actttgtagg gattttttgc acaaagttgt 240
 tcattaatgc tgattcttta tctgcaaaaa caaacagtga tcgggtgtgt gaattntccg 300
 tactcccat taaaaatcat tggagaacgg tgctattgat aagcgttgaa tcatcccact 360
 tggcattggc tcatcttctt tactaatat 389

<210> 1020
 <211> 291
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1020

cacagacaca cttagtgggt gacattaact attagtagtg gaggataata caaccagttt 60
 aatgcgtggg tggtttgacg ttggagaatt gattntgagt ctaaaattaa tattagatat 120

atttgtaa atctggtttt atgttgga aaattcgaa attaattcta ggtccataat 180
 tgattttgga ttgaaacaat attgagtagt atctgcccta gattcaaaaa tttgtattga 240
 attttatttc taacttgatt ttataattaa acattcagac ataaatcata t 291

<210> 1021
 <211> 383
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1021

agctaggtn tgggcaatag cccccacct gacgtccca aggtctcctg acccccgca 60
 catatctcca ggtaccactc tgtggtcaac gaataaaagc aggaagttca cccttctaca 120
 ctctctcacc tcaagcttgt aggattatgg ggtaccaccac acatgtggta ctaggtggcg 180
 gtcgggcaat ggtgcacaac aagttttcca catccacaat gcgcgcataa acccaccatc 240
 ccctgttgcc cacctccaac tgagctcag tactcccacg tagcccatat cctcgtttct 300
 ctcaacaccg ggtcccccac aatcctccca agcttccaca acatccaagc aaaacaacat 360
 tcanatagca caagctatca cag 383

<210> 1022
 <211> 396
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1022

ntatcanatg gatgtaaaga gttcattctt atatggctnt attcaagagg aagtatatgt 60
 agatcaacct cctagatttg agaattcaga caagcctaatt catgttttta aattaaaaaa 120
 aaggctttat atggcttaaa gcaagccccct agagcttggt atgagcgtct gagtaagttc 180
 cttttagaaa aggattttct tagaggcaag gtagatacta cccttttcat aaatagaaaa 240
 tcacatgaca tintactggg tcaaatttat gttgatgata ttcatttttag atctactaat 300
 gaattattat gcaaggaatt ctctcatgac atgcaaagtg agtttgaatt gtcaatgatg 360
 ggagaactct aattgtttct tggatacaaa ttaaac 396

<210> 1023

<211> 325
 <212> DNA
 <213> Glycine max

<400> 1023

agtgtttctt ttgcaagaag aagggacaca tgaaaaagaa ttgccccggg ttccacaaat 60
 ggcttgagaa gaaaggtgaa tcaatctcat tagtatgtta tgaatcta atgggtagtg 120
 gtaatattaa cacctggtgg attgattctg gatctactat tcatattgca aattctttac 180
 agggtagtga aaacctaagg aaaccagtgg gaagtgaaga aagcatttta tcaggcaata 240
 agctaggctc acatgtggag gccattggaa cttgcatttt gactttaagt agtggcttta 300
 ttttaaaatt agaaaggact tttta 325

<210> 1024
 <211> 381
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1024

taggagtntt ccaaccatca agttgacctt gcttgaagac ctccatcctt catggtatac 60
 ttgagagttt tccaaaaata tcagacaaga tggaatatgt aagggaactt catttgcagt 120
 aactgccat aagtatatct gtctatttgt ggagttgttc agttaccaag tagcactgtc 180
 atgttaccag aactgactga tattatagct nttggatgga aagggtagca atggctaaat 240
 cacgaagatg gtgaagaaaa agcgggttca atagtatctt caaaggtaga acggctntgt 300
 gcctaagat gcaaccattt attaattttt canaacggtt catgcacgtt gttcatgtga 360
 aagatttaga ctgattgata t 381

<210> 1025
 <211> 289
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1025

gcttatcttt ggttntacaa ccaaagtcca tgtgaacctt gagtaatcat ctactataac 60
 taagccatag taattttccac cttaaactcag ttctagaggg accaaataaa tcaatgtgaa 120

aaagttcaag gggttttgaa gtagaaacaa catttttact ttgaaaggag tttttaactt 180
gctttccttt nttacaagct tcacacaatt tatttttctc aaacttaagt tntggaagac 240
caattactaa gtctttccta actagatgat taagatgatg catatttat 289

<210> 1026
<211> 410
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 1026

ctcaagctta tgattggatn gataaatgtg aacgtgagag acctctgagt gatgcagaga 60
gtagctgacg aagagatatc gataggtgag gggagtttat ttttggttta tagctcttga 120
taccatactt ggngttagta aacccaacta tggggatgta tgattctccc tatgcatgct 180
agttttcaag aaaactgtgt ttntgactaa tgggatgca tatatttctt attgatgaat 240
gaaattgtga atgatattgt tgttttatga aacttgtgtt gtctgaagac ctgtgaaatg 300
tgaatcctan gcatgaaatt atatgtatat atgtggaatg cgattactga tgatgttaat 360
attgatgata atattgatat gaaatgatgt tgatattgag atgagatgat 410

<210> 1027
<211> 360
<212> DNA
<213> Glycine max
<400> 1027

aacaaaactt gtgctattca tctttttcat tctcttctcc ctttgccaaa aagaatttgc 60
caaggactaa ccacctgaat tctttttgtg tctctcttct cctttttcca aaagaacaaa 120
ggactaaccg cctgaattct tttgtgtctc ctttctccct tgtcaaagaa ttcaaaacga 180
cacagtctga gaattctttt gattcttccc tttcccataa acaaaagatt tcaaaggact 240
aaccgcatga gatattctttt gtttccctt cacaagttt caaaggacta accgcctgag 300
aactttgtct taacacattg gagggtagat cccttgtgga caagtagagg acatctactt 360

<210> 1028
<211> 409
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1028

tctatggagg ctggatcttt gagcttcaat gatgttnttc aatggtgatt ntccaccata 60
gagttgcagc ggaagataaa ggagaagagg tgagaggaga cgccatccac tatggaataa 120
gcatggaag gagaagcttc accaccaaga caatgtctta gataagaagc ttagatagga 180
agtttcaatg gaggaagaga atgagagaga gaaagtggca tggaaaattg aaggaagaaa 240
gggagagaag tttaactntg aagtgtgtct cacaagactc tcattcatca aagttgtgac 300
aagtgttaca catgtttcta tntatagcct angtcactaa catttcacgt gaatctaaga 360
ggaatattcc aagaatatcc canatgcac ttaacatatt ccaagaata 409

<210> 1029
<211> 521
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1029

ctttcggtcc gtctatngcg ngcccgcgat ccttagagnc gcctgctgca tgcaagcttg 60
angaganaag gngaatatcc tttttcttct tggatgaccc aangtggcaa cgtgcttcat 120
ttagttaaat atcgtaaca ggctcattca ttgtggacgg taaccacgagc gggaattcta 180
agacaatgat gactcacacc ttcaattcgt tggaagctct tctaacctaa atttgacata 240
ctacactgga agagcactta ttcatttgca ccganggggtg gcatatgctg ccatgatgaa 300
tcgaacatct agggaaccat gcccaactca cagaatttaa tatgcggcac actactaagc 360
ccaacgtgaa ttgcacagga aatcatggtt gcgcgcttta gaattgactt catcacaagg 420
gtagatacga taagacatgt actgtgtttg atacacctca agtccaatt gatcggcgag 480
tcatttagga gatgatcgtg aatgcagaca caacctcacc t 521

<210> 1030
<211> 87
<212> DNA
<213> Glycine max

<400> 1030

accttcaggc gactttctct aagcgccgca ccggaatctt caagaacgca agtgagcttg 60

ccaccctctg cgacgtggac cttgctg

87

<210> 1031
<211> 460
<212> DNA
<213> Glycine max

<400> 1031

aagcacctga gctgcagcta tgctgctata ttacaataga ccttctttac ctcagcagca 60
aatcaacca caacagcaca attatgacct cttcagcaac agatacaacc ctggatggag 120
gaatcacctt aatctcagat ggtctagccc tcagcaacaa caacagtagc ctgctccttc 180
cttccaaaat gttgctagcc caagcaaacc atacattcct ccaccaatcc aacaacagca 240
acagccccag aaacagccaa cagttgagac cctccacaa ccttccctca gaagaacttg 300
tgaggcaaat gactatgcag aacatgcagt gtcaacaaga gaccagagcc ttcattcaga 360
gcttaaccaa tcagatggga caattggcta cacaattgaa tcaacaacag tgccaaaatt 420
ctgacaagct gccttcttaa gctgtccaaa atccccaaaa 460

<210> 1032
<211> 419
<212> DNA
<213> Glycine max

<400> 1032

acaggcctat atgacatctc ggactatgat taactccctc taacctccaa gtaccagcaa 60
atccagaggt aactctacaa actctcaaag catcactctt tatcactcat agcactacat 120
tctcactatc taaccctagg ttaactctac cctacatctc tagcagattt ccataagcaa 180
ttgcaaaaca cagacatcac atgcatcatc atagacactt ctaaaccaga acgggaaagc 240
gtgactcaca cctgacatga cgaagttaac atgtttcagt gagattctga cagataccat 300
ccagaacata aacctagtgt actacccatg atatttccaa aacaatccca cagaatatgt 360
gagaagatgc taccaacctg aaattgaagt cccactatag ggcgcttacg actccgaaa 419

<210> 1033
<211> 448
<212> DNA
<213> Glycine max

<400> 1033

agctcgaact tgaaataggg tcaggattga tcttatcggt cctcatggct cttgaagaaa 60
taaagagaat gaaaagtaca aataaatatt attttatttg taaggattaa aaatacattt 120
aaacctaataa attaacacgg attatgacta tttttataa aattatattg ttgttttttt 180
tttaatttta gataacttat gattgcaata ttggttaatg aatataaact tcaccaaag 240
ttaatagttg attttatata attaaaacca aaagttaata ttttataata tatatatata 300
tatatatata tatatatata tatttcattt cagtaagaaa aaatcatatt atatataaaa 360
aaagctctat ataaatttgc atagaggggc tcaactctca agcaccttaa gtcagcctac 420
tagggtcac acaggtaccc gaagataa 448

<210> 1034

<211> 356

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1034

cgccagttnt tacgtatact ttggaagtat atatattaac atcacgcact tatatgcgag 60
taagagcagc tcttctacgt ggtcattgaa ggaagattta agtgttggta ttttttgta 120
acttacataa tgttgctctg atgatattcc gtgaagaatg ttaacgactt ctcgttttag 180
ccatectatt ggtcttttca ttcgccgtac atatatatga attatactta ttgcatgaga 240
atattcttta cttacttgag actattctaa tttgtgtgcc tttaccagaa tatcttcacg 300
ctacactgga atgacaatgg tggatgcctc tctatgagaa ataatgacca catgtg 356

<210> 1035

<211> 449

<212> DNA

<213> Glycine max

<400> 1035

agcttgtcat ggtatatata tgtttcttaa tagtctactc tgtgttatat aatacttagt 60
taaatatatt gtgtatacta aaagcaaag cttaacatta tacgtcttcc gtactcaagg 120
atataccaac attgaagggc ggactgtctt gatgtagcga ttttaacagc gatgacacta 180

tctttgttct atatattaac tgtcatggag atggagctgt ctttgtaaac aatctttgtc 240
 attatcagct gaaagatgaa catgttggtc gataggaaaa taatgaaata ctttatataa 300
 atgtattgta aagttttaag aataaactgc tcaaattgcc agcagtttgt cgtagaatag 360
 tagtgcaaat tcatgatgta aatgtattag aatacagaat aaacctatgg tgccggttatt 420
 aattggtatg agaacatctt gggctcttgc 449

<210> 1036
 <211> 82
 <212> DNA
 <213> Glycine max

<400> 1036

tagcacatgt tgttggtccaa tgaactcttc ttgacagagc atgtgttgaa caggaactct 60
 tagaatgatg tgtagaatga at 82

<210> 1037
 <211> 264
 <212> DNA
 <213> Glycine max

<400> 1037

gagcaacgca cagctcacat tctctgtttc aactcctaaa caatttttaa ttggtggctt 60
 ctggatttat gtcaatacat caaaatctta tgttatactt gtgtcatcat gtaatgcttc 120
 ctctactact gattcgataa aacagataaa aaacactaa aaaatgaaac ctaatatcat 180
 caacgacata aagcataaat tctagtatta gtatcaccaa aagttttggc tgctggtttt 240
 gtgcccattc ctcacatttg atct 264

<210> 1038
 <211> 429
 <212> DNA
 <213> Glycine max

<400> 1038

agcttctact tatgtggcag ggcgggcttc ctttctcttc ttgtctccaa cgcgaacttt 60
 gaccattggt cttccttccc gcgatgcttc ttttcatgtc tgccctgagt ggcttatagc 120
 ctaaaccata cttccacga ttaccttggg tatttatcag tctagtattg ccgccgttgt 180

tttttcctaa acccataaccg ggctcataac cggtccccaata cataactcgg gccatcatta 240
 ccgctgcac ggacagacta tgctgcccga agagggagtc cacggaggaa atgctgacca 300
 cctcaaaaga ctggaaagta gtttctaacg atttcttctgc ggcttccaca taaggcatgg 360
 aggatgggca gcttaccaag atatcttctc cgctgacac gatgaccaag tgcccctcta 420
 ctacgaatt 429

<210> 1039
 <211> 437
 <212> DNA
 <213> Glycine max

<400> 1039

agcttcgaag ggatgaaagg gatgtgttga gggatgatgag tatgatatta gaggataagt 60
 tgaaggcttg caaagggtcat aatagagttt gaccgagcag ttgagttata cagacgagaa 120
 catcttgaca atcattgatg ggtataaaga agagctaagc ctagctgcca gtcataaggta 180
 gagactagag gatgagcacg cgaaggtagt ggctctgcaa gcggaagggg aagcaagaga 240
 gagcgtgata taatcattgc acaggggaagc cgtgaaatgg atggatagat tcaactctcac 300
 tctgaatagg agtcaagaac ttccaaggct tttagccaga gccaaaggaaa tggcggatgt 360
 gtaccagct ctcgaggaag ttcattgggt tctcgattat tgccaacaca tgttcgaata 420
 gatgtgcctc ataatta 437

<210> 1040
 <211> 408
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1040

tcatgatgat gaaccaagca attttgatga tgccaaaagc ccttgatgatt gattcacgac 60
 ttcaagatca agcatcaaga atccaatcca agattcaaga ttcaagagaa gaaatcaaga 120
 agcaacaagt caagacttca cacatgataa gtattaaaag attttttcaa aaaccaaata 180
 gtatagtnt gttttacaaa agaattttct cacattgtct aagttaccaa agtgattact 240
 ctctggtaat cgattaccag ttatcagtaa tcgattacca gtgaccaact tggtttccaa 300
 aatgttttca aatggtttgc aacgttccaa aatgattttc aacagtgaat cgatacacta 360

tatattaatc gttacagtga atctgacgtt gaatcaatct attggaga

408

<210> 1041
<211> 423
<212> DNA
<213> Glycine max

<400> 1041

agcttctgaa gcatctacaa gatgtattaa ctgtgttagt tgtcagccat gattagaggt 60
tgcttcaata tactctatta gtttttgttt atctggctat tcagtcatta ttgatgctca 120
tttatcgtaa tatgtatggg tttgtctttc tcgtgattga tgatttgcca ataccattag 180
aaattctaac ttcgattgtt acgaaagact gatttaatgt gagcttcaaa attgtgctga 240
tatatagatg gataaacaca tgacgttgat acttatatga ctgtggtgct tcaatgtata 300
cacttggttt tatgtaggca acaacttcta gaatgtggag gctatgagct cgtgcacatc 360
atgaactcaa taatcactta tgtgataatg tatggctgga agtcagtaaa cggaatcttg 420
agc 423

<210> 1042
<211> 437
<212> DNA
<213> Glycine max

<400> 1042

tctcgccatt gacaatggcg gtacgcgtat ctcgccagta cttctggcga catccatggt 60
aaaacagacc cctctgttaa atacttataa aagagacccc ttacgtaaa tagtttgtaa 120
aggtgaccct ctacagtaaa ttaccactt taaaataacg tttggacatt ggattttcat 180
ataagttatg gatgcttata taaatagtgt gttggaagca atgcttgata acccattttt 240
atgcatctaa gttaagttac aagaatatac ctattacttc cttggacgca tgtttagctt 300
gcgtagttca ataaactata tcaattcacc aaagttaa atctcttatt atcattagaa 360
ctatgaaatc ttacgggct tcacattatt catgtttttg gttccggcaa aatgttttga 420
ccaacttgcc atgtaag 437

<210> 1043
<211> 407

<212> DNA
<213> Glycine max

<400> 1043

taacttgagc atctctgact atgaatcatg caacataata gggaacaaat tcatagtaac 60
cctcgagtac aagaccgaga tgactataga gtagtatccc ctcatatttg atgtcccat 120
tagcaacaga cttagcatat gttttatttt catttgaaaa tgttgtagaca tgtgtcgac 180
taaatccaac gaataaacac aaaatacatg ttataacaag gattctgtga taaaattatg 240
tgcacctcag gacgtaattc tataacatgt tcttagttga gtacgaggct ttacaccttt 300
tgcttgacaa tggcaaagga gatgcacata tagagtaact agcgagctat ctaaataacc 360
tctcgtgaca ttagaccagc ggaaagaaac atgatgacca aactctg 407

<210> 1044
<211> 325
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1044

ttcggccgac actggcgtgt tcccatgcac tccggcgaga acacattgac ccacctcgat 60
cagataaaaa gacattaacc accggtcttg atcggaaaaa atgctggttg acgtcggcca 120
tgatagatga ccgatcgagg tctgaaaata aaagaatcac cggatgacgc cgatcgagca 180
tatacctaatt gacatcatcc aaatattatc cagggatttg atagaaaaaa caatagctga 240
taccagtcgt tatgtagtcc cgactgacat ttgtcagccg acattgcaca gtattntttt 300
caaacgctgg ccgataatat atctt 325

<210> 1045
<211> 344
<212> DNA
<213> Glycine max

<400> 1045

gctttacgga cctatcaaac tcagctagga ggattttgga gacccgctgt tgcgatatac 60
gaggatatgg gctacgtgag agtacgtgag cttaattgga ggtgggcaac acgagatggt 120
gggtttatgc gcgcattgtg gatgtggaaa acttgctgtg caccatcttc cgaccgcat 180

ctagtaccac atgtgattgg taccgcataa tctacaatc ttgtgatgat gaattgttga 240
aagaggagac ttcctgcttt attgttgacc acatagtggg acctggagat ctgttgccga 300
ggtaggaga ccatggggac gtcattgtgg gtgctattgc ccaa 344

<210> 1046
<211> 420
<212> DNA
<213> Glycine max

<400> 1046

taatgacatt gattatgaca tcacacgact tgctatTTTT agtttcattt tttcccaaga 60
aatatcgtgt acctttcgta aaagaattct gttttcgtcc ttttgtaagg aaaaaaaaaa 120
aaagagattc tgattgaatt tgagtaaact attttctaaa ataattattat tatgagtgc 180
aacttttttc ttatcttaac actctgtttt gctgtatatt aagactctga ctcaaatca 240
tcagacttgt ctataaaata agtatcttct acttccatcc cagtaaaaat cccacatgaa 300
ggacttaaca aagctagatt actttgtgac tcttataaat ataactaaga tgattattac 360
gaacacatgc tatgggttta tcttcgaaaa ggaacacatg aaataactcg attttaattt 420

<210> 1047
<211> 376
<212> DNA
<213> Glycine max

<400> 1047

agttggaagc aaaacaaggg agcaagcttt gggaacactt tcttcaagaa ctaaaacaaa 60
gtcctttaac cttttccatt ttcattcctt ttactatcct catgtatttc tggattggat 120
tcttctcctt gcatcagtag ttctacaaaa tagaaggagc agaacacaag gaaattagat 180
tatttaggat agtcacatat ctcttatgtg ttttaatttaa gtagcttgaa ccattacgta 240
ctaactctga tcaactctata tgtctattgc ttgctatatg attcgctaac acttttgaca 300
aggaactgga tgagatgaag cacataacta ctggaatttg tcgcaagtta ttattacatc 360
attgtagttt atgttt 376

<210> 1048
<211> 448
<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1048

agctgtgctt ctacactacc agcaacaata tggtatcaaa gagaanaact ctagatgagg 60
gttcactgtt atcaagcaag tcggagaccc agcatgacca caaattcacc tccactcctt 120
atgttcccat ggacccgggt atagggcccc ttttcaactc accgtgtgta caaatagtgt 180
tggtgtttat gtgcatcaaa tgaataaata tctatctcat gcttacattt caaaagcaca 240
ctaaaagcaa aaaagagtta tatacaagaa cgtaaaggaa ataaaaggaa accgacaaaa 300
gaggaagtca tgatattgca cgagattaga aggcctaact ctctaaaaac agtccccagt 360
ggagtcgcca aatgtcgcaa cctacccttc ggcgaggagg cgacgcggng ctcacgggtg 420
tgtcttccaa gggaggaagg ctcacgga 448

<210> 1049

<211> 447

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1049

tgtaggcctt gaatcttctt catcaatgga gtcctttgct tcttgaagat aaattggaag 60
cggaatggag aaggaggaaa ggtgattaga gatgccactt caaggagaag atgagtcaag 120
aacaagttca ccgcatatg aagatatgga tataagagct taaaggtagc agaagatgag 180
tgaggaggaga gggagagaaa gggcacgaca tttatgcctc agatgaggta tgaaatgtga 240
agtgtaatct ctcnatgat caaagttaaa aatatgcaca cacaaggcct ctatttatag 300
tttaagtgtc atacaaaatt ggaggaaaat ctgaatttct attcaaattt cacttgaatt 360
tgaatttgtg gagccaaatt tggagccaaa atttcactaa ttaggattgc atcatccctt 420
ccnctctgaa aatgaattga cctcaaa 447

<210> 1050

<211> 448

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1050

agcttgtgca aatcaaatca ctctacgtc tcctctctag catgcatttt ctttctttac 60
 ccactcctca cgtgtggttt tttagggaaa aaacaccata actaaacgcg ccgcatggga 120
 tccctatcgc accagatcca aatctagaac gatgggtgat caagaggaga cacaggaaca 180
 gatgaaggcc gacatgtcgg ctctgaaaga acaaattggcc tccatgatgg aggcctatgtt 240
 aggtatgaag catatcatgg agaagaacgc ggccaccgcc gccgctgtca gttcggctgc 300
 cgaagcagac ccgactctct tagcaactac gcaccaacct ccctcaaaca tagtatgacg 360
 gngaagggac aactgnggc acgatggcag ccctcacctg tgatacaacc gagcggctta 420
 cccttatgga ttgccgccca actattca 448

<210> 1051
 <211> 399
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1051

ctgaggggaag aagccacaag aacacttata tgaaggtcgg atgtataaga agaacatgta 60
 ttgctctccc tcccttgaag aactcgtgca caacaatgga gaataaagggt tctaagtttg 120
 ttttttcttg gagaagcgag gacatataag gctttatgct tgcttcaaat gaaacttggt 180
 tacgcttaat gttgacaaga tcaaactgat gacatgaata atcatttgat agccataatg 240
 ctgccatata tgcatttctt gccttttgat tntttaacta gaaatgacta aagtcgactt 300
 aagcaaaaat ggtaaaaact ccttctgtaa aactgaaaac cttatctaata ctttagatag 360
 tgtgctacat ccttgatat gtgactcata ggaacttgc 399

<210> 1052
 <211> 448
 <212> DNA
 <213> Glycine max
 <400> 1052

agcttctccc tataacagct tcaaaattct atattcagca cactactgta atttcaattc 60
 actcatgtta ggtggcatcc cttaaattggg aatctatagc ctggccatgc ttaaattgctt 120
 ataattttct agatgttcca gcttaaattgt tttgactggt ctcagcatga gaatttactt 180

tcattttgtc gcctaataca atactaacat aaatttccac tgatcgcttg aattgtaata 420

tccgataatt tctgttaaaa tgaaatc 447

<210> 1055
<211> 446
<212> DNA
<213> Glycine max

<400> 1055

gctctctaag tgaaatcagg tgcagccatc tccctaagag tcctctcaaa aggtggaggt 60

tgagccatgt tctcagtatg aaaattagta gtcgaatgct caaattcaga atgttcagaa 120

tcaccatcaa cataatactc agaatgctta aaatgctcaa aatgcacaga atgatcagga 180

tgcacactat gcctaagtaa tccatgaaag gttctatcta tttcaggaag ggttctaaat 240

cacctggatt gccctagtc atgcattata tgcagcaaat catgtgtttc tcaaacaagc 300

accagtggag ggtaaaaact acaactatag tcaaatgata tccaaatgag ctgaaatttt 360

atgagtaaca ccctaaaatc atgaaaagat agaacaaaaa tttgcagact aaaattcact 420

aactatgaaa actgactaaa gaaagt 446

<210> 1056
<211> 204
<212> DNA
<213> Glycine max

<400> 1056

tgctgggtgga gcttcgatgt atgctgaatc tttgagcttc aatgaggtcc ttcaatggtg 60

agtgttcacc atggacacgc cacggaaagt cataagataa gaggataggg gaggcacctt 120

tcactatgga ataatccaag gaagaaggag cttcaccacc aataattgcc ttggataaaa 180

aacttgaca agattctttc ctgg 204

<210> 1057
<211> 369
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 1057

agcttctagt cgtgcataga ccttctctng ggtctgacta tcaaacgttg catctgtgca 60

ttcatcgcat tctaatacag acgttgagcg cegtccaact gatggtactc gtgaccacca 120
ccacctgcta cagccataat ttaacaggaa aaaaaaatgt gcaataaaaa ctattaaggt 180
ttcaagacct cacaacactc tactcacgtc tgtagatgt gagtacactc gtgtttaacg 240
ctctcaatat gctatcgtga aatgtattcc ctcttgccct ttaccactcg agtggactct 300
taagttcctg gatggaccaa attacacaca catggtaata tttaatcaga ggagagacta 360
tatgatgat 369

<210> 1058
<211> 441
<212> DNA
<213> Glycine max

<400> 1058

agcttgccctc aaagaggtcc aggattgata atgttgccga aggaactagt tccgctcccg 60
agtatgacag tcaccgcttt aggagcgctg tacatcagca gcgcttctaa gccatcaagg 120
gatggtcatt tctccgggag cgacgcgtcc agctcagga cgacgagtat actgatttcc 180
aggaggaaat aggtcgccgg cgggtgggcat cactagttac ccccatggcc aaatttgatc 240
cagacatagt cctcaaattt tatgtcaatg cttggccaac agaggagggc gtgcgtgaca 300
tgaggctctg tgtgaggggt gagttgatcc tgtttaatgc agatgctatc ggccagctcc 360
tgggatatcc gttagtgttg gaagagggcc aggagtgtga gtatggccag aggaggaatc 420
ggtctgatgg gttcgtatgag g 441

<210> 1059
<211> 407
<212> DNA
<213> Glycine max

<400> 1059

cgcatgataa atactgggac agtctcaaac cctgatgtat cagtttcaga tccgtggagc 60
caatgcgcag tgggacaagt agagtgccta aaatcattgg tcagactcct acctatgtgt 120
gcctcgggag tcttgatgat ggcgccccaa ggctcattct ctaccctgca agcaactacc 180
ttggaccgaa agctattttg caatttcaag atgcctgcag ggctcctcaa tcttatcatg 240
atattgacct tatcaataga cattcccttg tatgaccgca taatggtacc tctactagcc 300

aaatacaggg gcttgccgaa tggattctgt agtaaaactc caattgggat tggattgctg 360
 tttgtatgcg cagctaaagg aacatcagct gtagttgaaa ctattag 407

<210> 1060
 <211> 430
 <212> DNA
 <213> Glycine max
 <400> 1060

gttatctcgcg catattttgc tgggtgcgct ccatcatacg atccatgaca cgccatgcat 60
 cctatctgcg gaaaaacaca aaatgcttag cgtactaatc accgtagctt gttaacatga 120
 acgtattaat aaatctagta ctgcgctcac tcacctatga ttccggccct gagaagaaaa 180
 tgaatctgga aaatgagaag gcaacaacaa cagcgcgtga cgtaaactct tatgataagg 240
 ggagagaaat gagattagac gcttacgcta tatagaacga tgcattgccg ttcttttagag 300
 atgacgtgac aactaggtg acctcttttg caaaactaaa tttggggccc ttgtactagg 360
 tactatacct tacaacgggt tttcgggtgta atgttttcta agatttacag agagttatac 420
 atattgttct 430

<210> 1061
 <211> 380
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1061

atcaattcat attggatgca ctccaacttg taatgatgcc ctattccttt cctcagagac 60
 tgatgccatg acaccgtcac cctttctttc agaattcagt gaaaaccct ctctcttgtg 120
 acaactttcg cgccttataa cgacattcaa atgttctaaa aattaattgg agagtgaaga 180
 acaaacctca ctacgtgata attgttcccc catccgggcc ttgtgcaaac attcatgtga 240
 taagattttg aaagtgggaa tagtagatcg aggccaccta aatggaaatt gaaaaaaggg 300
 tgtcaggtac aattatcaca cactntttta caatgatata ttgcttagaa ttcaaaacat 360
 tcctaagtga cacaacaccg 380

<210> 1062

<211> 439
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 1062

agctntgttt catagataca tatatacaac taatataaaa tgtaacaaaa atgatgaatg 60
 aagataaaaa aaaatattgg aatgaaatgg ctatgacaat gatgaagatt aacaaataac 120
 cacacgaaat attcaacaca attcatatta aggatcaggt tatgaatttc aagactattc 180
 tcaaataaac atgacatgat ataacacatc agtattgctt atattagttt actttaaatg 240
 tagccttatg gtgaagtaag taggtccttt gctgtaagat ggaaggatta actagaagac 300
 acatggcatc tgtagatag taatgacagc atgcactatg ttatatgcaa caaagatctg 360
 gtgaacccaa ctatactggc tggatggaca aaactcggag atttctatgg actcacagga 420
 tatcatcaag tgaccatga 439

<210> 1063
 <211> 410
 <212> DNA
 <213> Glycine max

 <400> 1063

ctatattttc agtagatgaa tatgaatccg cggccacctc atgtactcct ctaaggacaa 60
 tagcatcatt tggtgcactg aattgttagg agttggaagc catcttctca atcaaactcc 120
 tagcctcagc acgggtcata tcaccaagag ctccccact agcagcatta atcatactcc 180
 tctccatggt gctaagtccc tcatagaaat attgaggaag gagttgctca gaaatctggc 240
 ggtgagggca gcttgcacac aatttcttga atctttacca gtactcatatc gagctctctc 300
 cactaagatg cctaattgcct gaaatgtctt ttctgatggc agtggtccta catgcaagga 360
 ataatttctg caagaacact cttaacgtcg tccaagctga aaatggacct 410

<210> 1064
 <211> 432
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 1064

agctatgctg atttggctt cgccagtga atgttcgaag tggatctgan aagaggcaaa 60
 tttaatcatc ctgcttagac gaatgagaaa actgctggcaa ataaagaggg tgaggatgag 120
 ggagaaaccc atgctgtgac tgccattcct atacggccaa gtttcccacc aaacccaaca 180
 atgtcattac tcagtcaata acaaaccacc tccttaccca ccaccagtt atccacaaag 240
 gccatcccta aatcaaccac aaagcctgtc taccgcactt ccaatgacga agaccacctt 300
 tagcacaac cataaaaaac accaaccaag aaatgaattt tgcagcgaaa agcctgtatg 360
 attcacccca tattccggtg tcatatgcta acttgctccc atatctactt gataacgcaa 420
 tggtagccat aa 432

<210> 1065
 <211> 422
 <212> DNA
 <213> Glycine max
 <400> 1065

tctatataat ctgaaccatt ctatcaataa acacacgtcg agttgtattc agaattattag 60
 agtttatctc ttattatctt agtgagagtg attctcctaa attcttgagt gattcaagaa 120
 caccttggct gtatcaaagg actttcacia cctttgtgtg ttgccctcgc tggaaagagt 180
 gattctttct ttactttcat catcaccctt gttctttcaa atcacaattc cagaagatcc 240
 acctctgccc agagatatct cgtggccata acttccattt tacgcactca aattaagaga 300
 ttcttgagcc tatattgaat ttcaaaacga gacccttcac ctggttatgg aatcacctca 360
 tttggagccc tgtagcttca gtattgcat ttctatattt ctgtccagcc accacttaac 420
 ct 422

<210> 1066
 <211> 435
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1066

agcttcataa atagactgaa tcaaggagaa aattatcgtc gacccccaac acataagtta 60
 tttaacacca cttgtcacgt tgttgactag ttaatataggt ttattatggt tgcataagaa 120
 aacatgtttc taaatgtatt aaaaagcttg ccttggaata atcaagttat gagtatggac 180

tagagatgag atgcaaactg caatcaacat aggggtacaat aacaaggctt gttctagatt 240
 tttttttttt ggaagttgaa taaagtttaa tctagttgtc tttttacagg tatgcttttag 300
 gtttacttag ataagacatg aaacagaaaag tgaccactct tgattaacta gatattgagg 360
 cccttgtaaa aggtactaga ccttacaaaag tttttcgcat ttaattttct ctaaatacta 420
 tagatatttg ataca 435

<210> 1067
 <211> 428
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1067

ntgatctagt gtaaaattaa ctttccttan aagataaaac attcacaggt ccagatgatt 60
 accgaatttg taagaacaac aactaagtca tcttgaaagt actcaaacag ataatagtgc 120
 actatagctt ccctcagaga cagaggccat gaaaccttca ccctttcttc caaaattcag 180
 tgaaaaatca gcattcaaaa gtgaaaactt ttggcatggg aaaggacatt aaaatgttaa 240
 aaaaattaat tgggtgaagga agaacaaacc tcaactagtga taattgttcc ccaatccggg 300
 ccttttgcaa acattcagct gataagattt tgaaagtggg aatagtagat tgaggccacc 360
 taaatggaaa ttgaaaaaag ggtgtcaggt acaattatca gacactnttt tacaatgata 420
 tattgctt 428

<210> 1068
 <211> 442
 <212> DNA
 <213> Glycine max

<400> 1068

agcatttaca atgattaaga tatactcttt caagtttttt ggccataaat tagtctggga 60
 tctcaatcaa gtctcgagac tcttgaaggt caatggtctt taaactcacg aggttctgta 120
 aaaaaataa aaatacatca cattgaccat aaatcattaa aataaaattc ttagtatata 180
 ttgaagctca tcatcctagg tgtggctctt gtgattgtga atgagtttca ttcgagtaag 240
 caaaagacta gttaccaatt tgtcttacga ttctctaccc tacctttgca aatagaaaga 300

aaaatgtagt atgtatattc acaatcatac ctgaacccca tcccagagct ttttgagctt 360
gctacgaggc atggaaatct ctacaagtcg ttcagcgcag aagttatacg gcaaagactc 420
aagacaacat tcatgccaat ga 442

<210> 1069
<211> 450
<212> DNA
<213> Glycine max

<400> 1069

gagctttatg gaggctggat ctttgagctt caatgaggtc cttcaatggt gatTTTTccac 60
catggagatg cagtagaagg caaaggagaa gaggagagga gaggcacat ccactatgga 120
ataagccatg gaagaaggag catcaccacc aagaatgtgc catggataag aagcttgaag 180
atgatgcttt aatggaggaa aagaaagaga gaagggggga gcacgaaatt gaaggaataa 240
aagagggaga gaagtggaac tttgaagtat gtctcataag acgtccattc atcaaagtta 300
caacaagtgt tacacatgct tctatttata gactaggtag cttccttgag aagctttctt 360
gagaaaactt ccttgagaag cttctttgag aaaacttctt tgagaagcta gagcttaact 420
acacacacac ttctaataac taagctcacc 450

<210> 1070
<211> 448
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1070

agcttctcnn cctatttgc ataaataggg ggatatgtga agataaaaaag ggttcagccc 60
cttaggcact tctctctctc tcgaaattgc tgaggaaaat tatttccgtg aagaagaaaa 120
gggttcagcc ccttaggcac ttctctctct ctcgaaattg ctgaggaaaa ttatttccgt 180
gaagaaaatc caagcagagg cgctttcgta acgtttccgt gagtaattac gcgaagattc 240
tcgaccgttc ttcaagattc atcgttcggt cttcgttttc ttcagtcttc aacgggtaag 300
tacctcaaac cgagcttttc aattcattct atgtaccgt ggtggtccac atnttgtttc 360
atgtatttnt attctcggtt tcatttgctt tttatacccc ctnttgacgt gcttaagcca 420
tttatntaag tcatttctcg cttaatct 448

<210> 1071
 <211> 107
 <212> DNA
 <213> Glycine max

<400> 1071

tgctgactgc aatcatcttc gggtagacattt acaaggctgg tagtagaaat cttgtaaatg 60
 gaagtcgagt caaccttgat gcagatgact ttatacacgc atgcttc 107

<210> 1072
 <211> 384
 <212> DNA
 <213> Glycine max

<400> 1072

tggatctgac tgccttgctt tatgtgatat acataattgg gtataaagac gactagggcg 60
 cagtaactga cgtttgcttt tatatccaaa ttgtataacc tgtgataaat catttaaagc 120
 cattggctct ttattattca ttatatgac aattctttta ccacttctag attgtgagta 180
 tcaatgatta ttcattgatca caaagcatat gccaaactca gatgatagta tatcaaatta 240
 ctatacttct gatcatgagg gcgagccctg gtgcatcggt aaagatgtgc ctggtgact 300
 tgttggtcat gggttcaaata ccagaaacag cctcttttga tatgcatggg taatgctgcg 360
 tacaacatcc ctaccccata cctt 384

<210> 1073
 <211> 318
 <212> DNA
 <213> Glycine max

<400> 1073

agctcctgcc tcagttaata ctcatccgtg gctgtgtgat cagcttttag aagatggcca 60
 cctaggattc ttgtgtgcat tttccacacg caagaggggt aaccgaaaca catccagcat 120
 ctaagtcggg aatcgaagtt catggaataa tggccgacat tctcaaattg gtctatacta 180
 gattgccttg gatgagttga aactgtcggg gcattatctg aaaatgtgcg tgccctgtaa 240
 aggaccatat ggataaagga gctgaccacg ccatacttat acagatgtac ttgtgctacc 300
 tacttttccc tgattacc 318

<210> 1074
 <211> 356
 <212> DNA
 <213> Glycine max

<400> 1074

agcttgaggt ctggaagcag tgcttttact gtttctggat gctatctaca ataaaaagaa 60
 cccatcagtt tattagacca gaagttatta agattaaaac agaaaataaa aacgaaaatt 120
 ggcgatgtgc gcttagcgag atgcagctag cttagcacgc cttagtaaaa acaacacacc 180
 ggcttagcgc aatatgggtg cgcttagcca gtcacgacaa agaaattttc tctgcataat 240
 tggctttgcg agcagtgcta gcttagcctt atgcatgccg caacgaatag tgcttagccc 300
 atggggatgg cacttatccc gagcaacact tccaaaaatt tgactatgta atctgg 356

<210> 1075
 <211> 252
 <212> DNA
 <213> Glycine max

<400> 1075

tgctagctag ggtaaatctc aaagcttcat aagcatatcc ttgtcagagg actgtcttta 60
 gtcacatcat caggatgatt ctctatgtga agcttgccct aaagggaaac aagtgaaaag 120
 ccattgctat aaccaaaaaa taccgtttgc acttctacgc ccttaacgcc tatgtggccc 180
 aactataact acatccctct ctggacacat atatgggttg gtcacgagg actattacac 240
 cagatggaca tg 252

<210> 1076
 <211> 412
 <212> DNA
 <213> Glycine max

<400> 1076

gttgcttcct ccagaaggca tcgccttctg gggaactacc tggaaggccc tagtgggcct 60
 ggtttctatt tgcacccctt attactaaa tacacccctt tacctttttt tgctgattct 120
 ttttccgtaa cgatacggaa ctttacgaat tacgtaacga tacttgtttt ctattcgtaa 180
 tgacacgaca ccttacggat tacgtaatca tcccttcttt atcttacgaa atgttatgat 240

actttacgga ttgcgcataa acactttctt ttgacttccg acatgtcacg aaacttcacg 300
gattgtgcaa cagtgcattc tttagacttc cagcatgtca cggaactgca cagattgcct 360
aacgatgtgt gctaactacc tacgagtggg catacgaggg tctcatccca ct 412

<210> 1077
<211> 382
<212> DNA
<213> Glycine max

<400> 1077

acacagacca ataccacaac tttccttact caaatacccc agtaacattg tcttcgttcc 60
aatttgttca ccgttggtac gactcgaaaa ttttactgga ggtccctagt acataagtct 120
acattttgac cgttgggtac tgctacaaaa cgtccataac ccaatatgta caaccctttc 180
cacaaccagc aatgcataag cattttctgc acaagcacia aattatgctg cacatttcaa 240
cagcaaaatt ctgcataata gtgcagattt tcgaaatcac tcttgccctc ttccaatggt 300
gcccaaattg gaccctacaa gtccctatata aagtataaat catacctaaa ccacagacaa 360
gcttcagacc aaagcaattc aa 382

<210> 1078
<211> 434
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1078

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agatatactt atattcagtt ctgtagatgt tgtttgcgct gatctcccat tagaattaga 120
gttctaccct gataatccat cttagtcctt taatgcagac tataattttgt agattaccaa 180
aatgaaact ctgattctaa ttgaataaca acatcaacag ccaacaatac caaacaacc 240
ttatctaagt tttattcttg aaaatatcac ccgtgatcac aagttcacag ctatgatatt 300
gccataattg ataaactgac ctgggataag aaatggctgg atattatctc atgcgatggt 360
aatgaaccag acaagttatt attccttact ccaccaagct aagaacggcc cactagtttg 420
atthttcaaca catg 434

[illegible]

agcttctttg	agaagctaga	tccttatcta	tccatacccc	tctattaact	aaattaattt	60
ccttaaaaat	aattacggat	gaaaataacg	caacaaataa	tcaaacatca	aacataatta	120
ctaataatat	atagatatat	atatcagggg	gttacatcag	cacctgcaca	acctaaggcg	180
cccgccccc	tccagagggg	ggctccccc	gctcgggctc	caaccaacgac	tcgttctgtc	240
ggcaatgcct	actttggatc	cggttccaat	gccatgagga	actttccccc	gaagccaact	300
ccagaattca	ccccactccc	aatgacgtac	aaggacctct	ttcgcgtccct	catcgccaac	360
caaatgggtcg	taataactcc	cggggaagatc	taccaacccc	ctttcccaaa	gtgggtatgat	420
cttaatgcaa	cttgcatgta					440

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<223>      unsure at all n locations
<400>      1080
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<210>	1081
<211>	436
<212>	DNA
<213>	Glycine max

<223> unsure at all n locations
<400> 1081

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cccgaacgaag aactgacaa aaacttatct tctccttctt ggacaaagta tggcgggctg 120
ggggaagta aattatcttc ccacagacc ttggatgcaa atgtgattgt atacccatat 180
cagctaaatc ttgacaggta ttcaagccat ccttcattctt gccttgaatg ttcatgagcg 240
ttccaaccac actgatacaa acatttttct ccacatgcat aacatcaata caatgtctaa 300
cgtcaagatt acacctgttc ggaagatcaa agataatgga ccttttcttg catatgcaac 360
tctgactttt atccttcttt tgagtacttc caaatgcagt atntaagtga tgaacccgct 420
gatatacctg ctcacc 436

<210> 1082
<211> 456
<212> DNA
<213> Glycine max

<400> 1082
tctgagtcac ctgccgatg caagctgaac ccaaccgatg agttgtgaac ttttactgtg 60
agtgaacgac tagctgtgac taataatctt tgcatcaatt tgtgaatttt agaataaat 120
ttataaatga ggacttgatg aaggccatga ttgtgcatat acaagccttt tgaccaaaaa 180
gcttaccttg aattataatt gtatcatttg caccctttgt gagctgaatg attttgtcaa 240
taattgaacc ctaaacctga atgattatct ccagatacct tgtttagatt ctaggagagc 300
atatgggttca aggcattacc ccaaatttag gggagtggaa ctaattggga tgcaaagaaa 360
gagataaagc atcagcacac aacaaataag ttgtgtgtta aaaaagaâââ aagacaaaga 420
aagcaatcga aggaaatgtg tgttgatgta ataagg 456

<210> 1083
<211> 432
<212> DNA
<213> Glycine max

<400> 1083
agcttatgag catgagggtt agccttttct tgactataaa aataacgttg tttagttccc 60

ttgcttactc ccataatacc aggagccttg tgcataatgc tttttataca taagctaagt 120
 agtttctggt gtaacaaaag tgtttgact acactatttg tcactacatt cgcggtacgg 180
 gctaaggctg attcatttat gcaggttgtc attgggtcaag ctgctgtgca cagggtgtaat 240
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 gacttgaca agctatctgt tgatgtagct ggcaatttgc ttggcaccaa taatgttaat 360
 attgacaaga cacctatcat ggccgtgaa gactatgcat tctatcaaga ggtcacacct 420
 ggctacttca tc 432

<210> 1084
 <211> 433
 <212> DNA
 <213> Glycine max

<400> 1084

agcttcggaa gatagtgatg aggtacattc cctataggca gagcttgaaa gagcctgcgt 60
 agtcgaagag aagttcaagt ccatagccat caaagtctga aaagagtatg atgaactaag 120
 ggatgtcaat atggccaccg atgaagcctt ggaatgagaa accaagaagg cccgaaagga 180
 agaacacgac caaagcaaag ttttgagggg ctttataggg cagcaatagt gagctcaaac 240
 tccgaagagg tgaaaagaat catcacgggt caaaggcatg atctggaagg acgagctaaa 300
 agcttgccct aggtcgaaaa gaaatttgtc ccaacagtta aagtgagact gaagggaata 360
 tgtgggcat catcgatgag tgcaaagaga agctaaatct agcggcaact cacgagcaaa 420
 ggctatagga tga 433

<210> 1085
 <211> 428
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1085

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 ctttcacttc ttttcctttt agaatccctt tcatgattat gtatctcttc taagaaatct 120
 gaaacatcat ctaacacatt ctttcttgga gaaataacat tagactcatt aaaggaaaca 180
 tgaatggatt cttcaatatt catagttctc ttattgtata ttctatatgc ttactatgc 240

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aaggaataac caaggaagat tccttcattt gccttggcat caaactttcc taagttttct 300
tttccattat ttaatacaaa acatttgcaa ccaaagacat gtaaattgtga aatgtttggt 360
tttctaccat tgaataattc gtaagggagt ttctttaaga tgggtcttat taaagcctta 420
tttaagat 428
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cacgacaaat aagttgtgtg ttaaaaaaaaa agcaatcaga gaaaatgtgt gctgggtgtaa 420
taagggtcaa agcaaataaa agtgaaaagc tagtgc 455

<210>	1088
<211>	422
<212>	DNA
<213>	Glycine max
<400>	1088

agcttcaaca	ttcaatgtca	agcgtctcga	tatattacgg	gactcaatca	tacatccgag	60
taaatagtta	ttgtcgtttg	aattgggtca	gagcttcaac	attcaatttc	gaggggtctcg	120
atatattacg	ggactcaatc	cgacatccga	gaaaaaaatt	attgtcgttt	gaattgggtc	180
agagggttcaa	cattcaatta	tgagcgtctc	gatatgttac	gggactcaat	cagacatccg	240
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cgatatatta	cgggactcaa	tcagacatcc	gagtaaatag	ttattgtcgt	ttgaactggc	360
tcagagggttc	aacattcaat	ttcgagcgtc	tcgttatatt	acgggactca	atcagacatc	420
cg						422

<210>	1089
<211>	406
<212>	DNA
<213>	Glycine max
<400>	1089

agcttcttat	ccaatgctca	tcttggtggt	gaatttcctt	ctattctggt	attagaacga	60
gtggatggcg	cgtcctatgg	aatataatac	tacttattgg	gctgctgttt	ctgagaggaa	120
tggcaccata	aagtgacctc	attatggctc	aaggattcca	ccttttggtc	ttccacaaag	180
ctgatgctta	agctgacaat	tgtgaactga	atgcaatttg	agccaatgtc	ctttatgggtg	240
actaacccaa	tacttgtatt	aggtgtatca	atttatactt	gattgcgctc	ttatttagac	300
agactaaaact	ctttaaaatt	gtaataaaga	ggtgtgatat	actcctacct	tagggccatg	360
atacaacggc	atatcttggg	cttttctact	tataacatgg	gcttgt		406

<210>	1090
<211>	464

<212> DNA
 <213> Glycine max
 <400> 1090

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 ccatcacgat tatcgtctcc ctttccatta ttgggggtac cacctgagcc gccagatccc 120
 tccacctttt gggcgtgttc tttgaatgat ccgtccccct ttttgcacat gttctgtagt 180
 tgcacacctat ccggaacctat atcacaattg tactgatact gcctaacaaa ggcaaccatt 240
 aggtccttcc aagaatggac tccggaaggt tccaagttag tgtaccaggt aacagctacc 300
 ccagtaagac tttcttggaa ggaatgtatt agcaattcct catcttttgc gtattccccc 360
 atcttctgac aatacatctt tagatgggtc ttgggacaag tagtccccct gtacttgtca 420
 aagtccagca ccttgaactt gggaatgacc atgtttgggt atta 464

<210> 1091
 <211> 449
 <212> DNA
 <213> Glycine max
 <400> 1091

ctgcagcttg caagtttaaat aaagaaactc ttaaattctct cttttatctt tagatttaaa 60
 catgttctta gttacattca attgtgcac atcgattcgt tttgttattc cttagcgaaca 120
 ttttgtttta tattgatgat tcacaactgt gtaggtaatt gtcgttgata aagttttaac 180
 cttcattttt cacatccctt atatttctag tcttacaaaa tgtcatttct cttgtatcta 240
 ttaattaaaa caaactttta ttaaaaaatt attatgagcg tgtttcgatc cgcaccaaag 300
 gcccttggac gtgcgtatatt catgggtcaaa catgagaaag tcagttgacc gtgatgtttg 360
 gttctccac tatactcgat tcacctcgtc gaaatcgatt ttgaagcaca gcatggttga 420
 ggcagcttcc acgtcaagtt aaaattgat 449

<210> 1092
 <211> 436
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1092

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 cttttaattg aaagtttga acgatgctta gtgttgaatt ctggcaatgc taaggtggtc 120
 ctctgaacat ggacaaatta tctgttaatg gaaattctgg ctcttcaag ggtgggtgga 180
 gggttctgat actgcctttt gagtagtggg ttttgttggt gccctttaga gtaaactctc 240
 acaatttcac aatccgattc tacaaataaa gtttatggaa cttccacgat tctacgtaaa 300
 atcgagagtt taacaaccat gattggaaagt atccgcgctc tctatctctt tatatacccc 360
 attaataaag ataataattg ttgatcagca gactgcctgc gttaagtgtg attactgctg 420
 gaaatgcaga tattat 436

<210> 1093
 <211> 425
 <212> DNA
 <213> Glycine max

<400> 1093

atttatttaa acctcccact gggcattaag ttaggcctac ttggttaaaa ggaaaaatat 60
 atattttttc atagaatggg gaacaactaa ctcaaattaa ttacaatga atatttaaaa 120
 atgtggagtg actgagttaa aatggattca cttaatcatg ctatatcgag tgactgagtt 180
 aaaaatttag agctatatat gtcgctagac gcttatcaac agtatattta tctggtttat 240
 aaattaaagt aaacgacaag aaattacttc ttatacattg tgctacgtg caatcacaac 300
 gatacaaaat tccatattat aagaattgag agtataacta tgttttcatt gagcatttat 360
 ttattaatgg tattgtgcta cggcatgtga ggctgtgca caagttcatc agatataaat 420
 attgt 425

<210> 1094
 <211> 192
 <212> DNA
 <213> Glycine max

<400> 1094

tgaagctcca tagctacgat tgacgccaat gactgtaact aacatgacta ccatactgga 60
 caggactgcc tccggatgga attgaggcta tctaatacat acaccagatt gaactcatgc 120
 aagctgttca cttgtcgcgc atgtagaggc tgctgtagac gcagtgaacg atgattcttg 180

actaaatgat ct

192

<210> 1095
<211> 429
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1095

agcttganga ttatggggta cccgtcatat gtggtattag gtggcgatcg ggcgatggtg 60
aaaatcaact atcccacatc cacgaatcaa acatgaactc accatcccca gttgcccacc 120
ttcaactaag ctcacgtact cctacgtagc ccttatectc gttcctttca acaccgggtc 180
ctcatcaacc cctccaagct tccacaatat ccaagcgatt caatttccaa atatcatgaa 240
ctatcctaaa ccaagaaaac agggcagagg cagaaaactc tgcccaaac acattcacat 300
attacaactg tccttactca aagaccccag taacattctc ttcgttcggg tacgttaacc 360
ataggatcaa attgaaagt ttactggagg ttcttagtac ataaatctac attgtgaccg 420
ttgggatct 429

<210> 1096
<211> 334
<212> DNA
<213> Glycine max

<400> 1096

agagatctac aaacattggt gtgcctacaa cactactact aacgaggaat gtggtagtgt 60
attacaccga gtttgatgag tactaggagg aactcgagag aagactctgg gatgagaagt 120
taactgattt tgcagacgat cgcatagaca ttgctattat gaaggaatct tacgcccaacc 180
tctatgacct caagggtaaa tcaattattc tgggtgaaggt gagaggacat ctaacgaagt 240
ttgatgaaga ctgcttgaa acattgtagg agaccccgat gactatggaa gaggggggaga 300
atgtgtgtgc tgattccaag tttgcactcc tgag 334

<210> 1097
<211> 439
<212> DNA
<213> Glycine max

<223> unsure at all n locations

<400> 1097

agctntacag cagatnttag taatgaccca cttttctaga attaaaataa cttaatgcc 60
 ttaacctagg gaattaaaac aaactaaatg gctgagtgt actgaaattg ttggcaacca 120
 aaagtcaccc ccaacagcca acaagtcagc caccatttgg tctcccaaaa ggctgatgcc 180
 taggttgcca attgggccct tattacaact tgaactaaag cccttttagt tgattaaccc 240
 aaaacatatt tttggtcagc caactttaca aggattgggc cattatttag acaaactaaa 300
 cactctaaaa ttgaaataaa gtgggtgcat ttagtcctcc atttgggcca tgatacaact 360
 cacaaccttg gacttttctc cttgaaactt gggcttgat tcaaatagta tggacagcac 420
 ttgttgaaga gcgtccttg 439

<210> 1098

<211> 380

<212> DNA

<213> Glycine max

<400> 1098

aggctctgag caaattcaaa cgacaataac ttttgactca aatgaccgct tgagtcccg 60
 agtacatcga gatgctcgta atagaaaagg gaagctctga gaaaatgaaa cgaccattac 120
 ttataactac gatgtcggat agagccccgg aaaatattga gacgctcaac attgaaaaca 180
 gaagctctta ggatattcca acgacaatat agtttgactc ggatgttcga ctgtgtcccg 240
 taatatatcg agactctcgc aaatgacaag agaagctctg cggaaattcg aacgacaata 300
 acttctgact ctagtttccg cttgtgtccc gtaatatatc gagaggctcg ttatagaaaa 360
 gggaagctct atgataaaact 380

<210> 1099

<211> 149

<212> DNA

<213> Glycine max

<400> 1099

catgcgtgct gggattgatc tgatgcctgc cttactaacg ccatagacgt actcttatcg 60
 acatggctcc ccatcgcaact caagtgcata acatgggaaa taaagagaga gatagctcta 120
 tacactacgg actatggcgt agagacctg 149

<210> 1100
 <211> 376
 <212> DNA
 <213> Glycine max

<400> 1100

catgagctag tgaacactct ctctgacatc atgtctgtgc tcatataagt gatcatgacg 60
 tgcattgaga tgttctctgct caactcgac gtcgtgcact gtgatacctc gctcctgaag 120
 acgcttcttt ctgatcttcg atggcactac actcttgagg ggaacattct gaagaactgc 180
 ctaccttgtc ttcattgttc ctctgacgct ggttacgac tttggagagg ttggacatac 240
 ctctcctgaa gatatgatac gcattgtacc tcactttag acaggggaacc aagtcacaga 300
 caccctcca tgetagecca gagttgggcc caattagcct ttcattttct acgcacgagc 360
 agtgaccttg tggcgg 376

<210> 1101
 <211> 438
 <212> DNA
 <213> Glycine max

<400> 1101

agcttatcga gaaaagaaat tgtataatgt ttgtttacaa cattgttaag ttcaactaaa 60
 accctttgta gagcattatt cccaagtgc gtaagaccaa ctgtaaaaga aaaaaatta 120
 aacacttgac aatggatgca tgcactacta tcactatacc agctagcttc attcgtctct 180
 ttcaagcatc tatagcaatt ctttgcaata aaatcttgaa actaacactt ggacagctag 240
 atctaaccgt tgttgctgga gtgtgaccaa attaattggt atatttatta tgaataattg 300
 aatattaaaa tactcttggc agtgcatacc taaaagctc acttgtggac aaaaacatta 360
 cgggtcttaa tggatagata agaattaaaa tcaatcaaag taaagatcag ggaggatcat 420
 catcaatttt cagcacc 438

<210> 1102
 <211> 447
 <212> DNA
 <213> Glycine max

<400> 1102

[illegible]

<400> 1103

<210>	1104
<211>	429
<212>	DNA
<213>	Glycine max

tcaagaatcn gatntcacag aatcaagatt caagaatgat cttttttcaa gattcaatca 60
agcttcatga atcaagattc aagagcaatc aagatcaaga ttcaggaatc aagaaaagac 120
tcaatcaaga taagtactaa aagtttttcg taacattgag tggcacaaga atttttcaca 180

aaatctttaa cgagagagtt ctactttctg gtaatcgatt accgagagcc aacattgggt 240
 ttcaaaactg atttaciaaag cttgtaatcg attaccatga gcatgtaatc gattaccaat 300
 attgtaaaat gttagatttc aaatctcaag agtcacaact agtgataaaa cattgtcaaa 360
 tcattgtaaa cttgtctaata cgattacaca atacttgtaa tcgattacca gagtttctaa 420
 acggtttga 429

<210> 1105
 <211> 430
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1105

agctngcttg tggggcttct atggaggctg gatcttcaag cttcagtga gtcctttaat 60
 ggtgattttc caccatggag atggagcaga agacaaagga gaagaggtga gagaaggcac 120
 tatccactag ggaataagcc atggaagaag gagcttcacc accaagatga gcctaggata 180
 agaagcttgg agaggatgct tcaatggagg aaaagaaaga aagagagaaa gaaagagggg 240
 gagcacgaaa ttgaaggaag aaaaaggag agaagttgaa ctttgagttg tgtctcacia 300
 gactctcatt tatcaaagtt acaacaagtg ttacacatgc ttctatttat agactaggta 360
 gcttccttga gaagctagag cttagctaca cacaccctc tcataactaa gctcacctcc 420
 ttgagaagct 430

<210> 1106
 <211> 429
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1106

ntaacctcat cgtctctcac agtctttaga tttgggagcc aatccaatcc ttgtgttcgg 60
 actctcagcc acttatgata gccgccgatg atcccattac ggcttccct aagctctctg 120
 tcctttcttc acgccgcatc ccatgccttg cgaactcctt ggagtaccct cgcgttgttg 180
 tcaactgaaac cccgtgcgat gaaaggcgtg atgctttcgt ctgatggcac tcctctcatg 240
 aggtagccaa gctgtcttat ggcgaggacg ggattataat taatacaacc ccttggtccc 300

atcaagggaa catttggaca tccttcgcat gaagatagaa tcctgattct tccttccttc 360
tagcgagggg accaattaac agacgcccct ccattgctagc caagagttgg tcccaattcg 420
cctttcctt 429

<210> 1107
<211> 438
<212> DNA
<213> Glycine max

<400> 1107

agcttgtcct tggtttaaac atgattggta catgatttgg gacttgtatg tattaatttg 60
ggaaaaattg gatgggggaa agactggttt tcgaaatctg cactttatgc agaattttgc 120
tgttgaaatg tgcagcagaa ttttgtataa gtgcagaaaa atgcttgtgt atggctgggtt 180
gtaaaaaggg tagtacatat ggggttctgg acatttgcta gcagatccca acggtcaaaa 240
tttacaccta tgtactagag acttccggtg aaattttaga gtcgatccga cggttaacga 300
attggaacga agaaaatgtt actagagtat ttgtatgtga aaagctgtga ttttgagttg 360
tgctttgggc agagtgtctg cctttgccct gttctgcttg gttgtgttag tacatgatga 420
tgggatgtgg aattacct 438

<210> 1108
<211> 469
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1108

ctaagctctg ctgcaatatt acaatagacc tcctcaacct cagcagctaa atcaaccacg 60
gtagagcaat tatgacctct ccagcaacag atacaaccct ggatggagga atcacccctaa 120
cctcagatgg tccagccctc agcaacaaca gcagcagcct gtccttctct tccaaaatgc 180
tactggccca agcagaccat acattccttc accaatctca caacagcaac aacctcagaa 240
acaaccaaca gttgaggccc ctccataacc ttccctcgaa gaacttgtga ggcaaatgac 300
tatgcagaac atgcagtttc agcaagagac cagagcctcc attcagagct taaccaatca 360
gatgggacaa tnggctaccc aattgaatca acaacagtcc cagaattctg aatagctggc 420

cttctcaagc tgtccaaaat cccaaatatg tcagtgccat atcattgag

469

<210> 1109
<211> 585
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1109

ctgtacatcc ttactngact ctaacctcac tattccctca atactaataa ctctcacacc 60
actcaccatc cgacgctgac acattganga ccgtcgaact gtagccgcg actctataca 120
tgctactgcg ngcatgctag catgcttgag gtgctgttat ggacgctgta tttacaagct 180
tcaatgaggt cctttaatgg cgaatctcca ccacggagat ggagcacaag acaaaggaca 240
agaggcgaga gaacgctcta tgcactaggg aataagccat gtgaagaagg agcttcacca 300
ctcagacgag cctacgataa taatctcgga caggatgcta caatggagga caacaaagaa 360
cgatagatag aaagaggggg agcaccaa at tgtaagaaga aaaagggaga gaagttgaac 420
ttcgacgtgc gcctcacatg actctcattt ataaaagtac cactgaggct acacacgcta 480
ttatatatat actacgaacg cttcttgaga ggcatagacc taagatacac atcacgcttc 540
ttatagcgaa gcgcacctcc ctcgagaagc tcccttaaca cgacg 585

<210> 1110
<211> 368
<212> DNA
<213> Glycine max

<400> 1110

ccgccgatga tcccattacg gcttcccta agctctctgt cctttcttca cgccgcatcc 60
catgccttgc gaactccttg gactaccctc gcgttggtgt cactgaaacc ccgtgcgatg 120
aaaggcgtga tgctttcgtc tgatggcact cctctcatga ggtagccaag ctgtcttatg 180
gcgaggacgg gattataatt aatacaacc cttgttccca tcaagggaa acattggacat 240
ccttcgcatg aagatagaat cctgaatctt ccttcttcta gcgagggacc cattaacaga 300
cgccctcct gctagccaga gttggtccca tttcgcttt cttttcgacc acacggtgac 360
cttgagcg 368

<210> 1111
 <211> 459
 <212> DNA
 <213> Glycine max

<400> 1111

ctgattgtat tttagagtga gtatttatgt caactacagt cttgctaggg tcaccacaac 60
 caatatatca acctaaagtt tatgagcaca gtcattgtag attttgtacg gacatgttat 120
 tcacttgtca aaattagtc taattgaaga tagtaaataag gaaaagttgt aaggactacg 180
 aatgtggtaa tcagcgtcgt caaaattatg gagcatgtga agttaatgtc ttgagaatga 240
 ttactttttc ttaagactga gagaagctag atcagttttt tgaatttgac ttctctatta 300
 tcccttttca ttctgtccca tgaaaataac attgcatgac atgacaatta tattaatatt 360
 taattataac atttaattct aaaaatcatc aaaactgtat aattttaaaa ggacaacaca 420
 aggacaagga caaggacagt acaaggaggg acaatagac 459

<210> 1112
 <211> 185
 <212> DNA
 <213> Glycine max

<400> 1112

acaaaagagg atcttttgca tatctatctt atgtaccgga agtccacat tcattatgac 60
 tgaattatcg ctgatcacat gttggctacc cccatgatcg tgtacctaat atgtgcctta 120
 gccatattta acactacgat aactgactac tacataattg ataccataa aacctttcca 180
 gctgt 185

<210> 1113
 <211> 612
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1113

tctctctcta actaccactc aacactcgtc acataattct agtttatatt gctataacag 60
 atacctgtgt atcgcatact cgcactctaa ctgcgcaact atctcacatc cctacaccat 120
 taçacatctt aatcactcta tatctgttcc acctactaca ctcgataaca catatttctt 180

cttattaaac ctatacatct acaatctcta tgnaactgcg aatcaccccta acctcagatg 240
 gtccagccct taataacaac atcatcagtc ctgctcactt gctttccaaa catgctacct 300
 aggcccaaca caagactcat tacattcctt cctaccaatc acaacaacta gcaactaact 360
 ctctagaaac aacacaacac attgaggccc ctctataacc ttccttcgaa gaacttgtga 420
 ggcaaagtac tatgcacaac atgcagtttc atcacgatac caatacctcc attcatagct 480
 taaacaancc gatgggacaa ttggctaccc aattgaatca acaacagtcc cagaattctg 540
 aatagctggt cttcttaagc tcgccaaaat ccataaatgt catgccatat cattgacgtc 600
 gctacacaat ct 612

<210> 1114
 <211> 394
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1114

tgtgtaagtt attatcattn gaatttctca cgagcttccg ttgttcaatt tcgagcctct 60
 cgacatatta tgcgcccga a tcggacatac gtgtgaaaag ntatgaatat ttgaatttct 120
 cgagagtttc cgatgtttta tttctagcgt attgatattt ataagcttga atcggatata 180
 cgtgtgataa gctatgacca tttgaattta tcagagcttt ggtgttcaat tctaattctct 240
 ctcatataag cgcccgatcg cgcattcgtg tgaatgttat accatttgaa ttacttagag 300
 ttacgatgct taattcagcg cttgtttata tttccttgat atccttcttg tataagttga 360
 cctctctgcc tgcacatctc tgtgatataa tagg 394

<210> 1115
 <211> 372
 <212> DNA
 <213> Glycine max
 <400> 1115

aaagctctcg agaaattcaa atggtcataa ctgttcacac tgacgtccga ttcaggctta 60
 taatatattg atatgctcaa aaataaacat cggaagttct cgagatattc aaatggtcac 120
 aatctttcac atggatgtcc gattcgggag cataatatgt cgagaggctc aaaattgaac 180
 aacggaaggt cttgagaaat tcaaatggtc ataacttttc acacgagatg tcgattaacg 240

cttataatat atcgatacgc tcgaaattga acaacggaac tcttccaaaa tatagatggt 300
cataacaatt accatgatag accattctgg gcctctatct tcatcagggtc gaataaacia 360
cctagctctt cc 372

<210> 1116
<211> 438
<212> DNA
<213> Glycine max

<400> 1116

agcttatata ccaccagcat cgttgtaata gggctgttga tggaacctct ccaaagtcaa 60
gctttccgca tgacttacgg aaagatctta gagttgacct tagcagaggt atccatagaa 120
accattgcat cactcaccca atactacgac cagcctttga gatgcttcac attcggagac 180
ttgcaattag taccaacct tgaagaattt gaggaacttc taggatgtcc tctcggggga 240
agaaagccat atctttcatc cgggtgtctc cctcttttga gcataattgc aactgtgggtc 300
taggatacaa caagaggttt ggaccgcata aaacagactc ggaacggcat agcgggccta 360
ccacggatgt acctagaaga caaggcgagg ggtatggcca atcaaggaga ttgggtcccg 420
tctatggata gtgtagct 438

<210> 1117
<211> 368
<212> DNA
<213> Glycine max

<400> 1117

atatgcgcat acttccttac atacgttgtg tagcacaaga cattatatta accgtaaaat 60
ataatgcccc catctacgat caaggcagct ccgacaccta aattatttac acgtacttgc 120
aacgtgtaga tgatacttac atcgcacaca tgtccttggc taaattgaca tacaagcata 180
ctcaaaacat tttggggtag gcaaaattgc aactgtgca cattatggca tttcttaaac 240
ctagacatac actaactcaa tgatgaatct tgactatcta cacaatacgg tgctacatgt 300
catgctcttt tcacatttgt gctccctaac accgcatgca aattcaagta tatcatcctt 360
tgctgact 368

<210> 1118
 <211> 494
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 1118

cgcgctgacc ttgtgaccct gtgtgcgca gcttaagaca acccagcatg ctgcttgttt 60
 gataaagaac atactacatt gtcgttgagc tgaagagacc cgcattacgt aagcctgtgt 120
 cttatatgac acacatatta gcttgactta tatatggcaa gggttggcgc atgaacatat 180
 atatgaacaa tggatatgac tgctacgtac gcctctctat ttcattggact tgacatcata 240
 cgagttcttc tctccccctt cttgcgtcta agatagctaa cgaatatcac tgacttatgc 300
 actgcgacaa gaactgccat acactactga actatcacia ctctgagtcc aacatatacc 360
 ttgggttaaca ttattatgaa cttctcaagc cagggagaac cttgagcaag ataccgagta 420
 ccgggatgac aatgacaagg atctgagatc catctttact ggttctatac tgtcatgatc 480
 actacgcacg ttan 494

<210> 1119
 <211> 222
 <212> DNA
 <213> Glycine max

<400> 1119

 tctcaatatc tgttcttgag tctttaacgc gctctgaccg gttattgaag ccgtgcttgt 60
 cgcttgaag agtgataaaa gaataccac cgagcatatg tggcgcacac tcatttactc 120
 actttacaaa cgaactctgg ccgatcgtgc acgctataac ctacgctaaa ccgctgagag 180
 gaaagtatgc aatgtatctt gtaacatggt tggttaactga tc 222

<210> 1120
 <211> 423
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1120

gtatgatacc tggttaacttt atgccacccc cggatttcga cattctcaac ttgaaacagg 60
 gtttaggggt gggggaacat aacaaaagcg tttaatcgga atgcattgat agtttaagta 120

ttttataatg aaatgcaata tataactaata tagtcgcttt tcctagcgat tcttctaata 180
catatatattg agatgattat gtaaaaatca ttatattaaa ttagtaatgt atcaaaacta 240
aaatttctaaa tatatgttga ggcatgactt aatttatgtt attttatcaa aataaactct 300
aaaatttatt ttaagaagct ttaagggtcaa cactataata taaactatnt agtgatacta 360
aactcgctca ttcattgatta ttgtcgcggt tacgaattca cttttactat taactcaaaa 420
agt 423

<210> 1121
<211> 412
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1121

ttcatgggct aagtaatata aaaagctctc tctgatatag taccagattg cttacattgt 60
aaagcttcat acgatacata ctanagggtac ccaactttgg ccatatgggc caatattact 120
aaacaataga aagggattta taaagttatt tctcagaact gaatagatta gagctaaacg 180
actatctatt acctgagata tgttgatgac aattacgtct acacaaattt tgaatatatg 240
ttcttctatt gttgaaatca attaaaaatt ataaaaattt gtcaatatta ttgcttctta 300
attaatactt tctataatcc tataattgat aatatgtgcg aactaactaa taaaacagaa 360
tgttttataa tatactacca tgttagaaat acttatatac tatcagtttc at 412

<210> 1122
<211> 392
<212> DNA
<213> Glycine max
<400> 1122

agctagacca atcctgaccc aaccgaggca tagttaatca gtgataacct gtgatgtacc 60
taaacaggcg agctcctggc agtcaaccga taagggaac aaagaccaca aagcatggag 120
gcttgtgtgg aggttggcca gctgcgaaac ttgtattgat atatgggata tggactctgg 180
taatcgatta ccaaggggag taatcgatta caaggcttaa aaagtaagac aggagactaa 240
gatgggtctct ggtaatcgat taccaggctt gaaaactaga tcatgaagct tggagggctt 300

ctcgtaatcg attaccatgg cgtgtaatcg actaccaggc ttataaatga gacttgaatg 360
 ttgaaggagc ctctggaaat cgataccaag ct 392

<210> 1123
 <211> 379
 <212> DNA
 <213> Glycine max
 <400> 1123

tggtgaggta gccatggaaa agcagagcgt ttggaatgat ttcgtaaatt tcagaaggct 60
 attgtgaaat gctggtaaaa acacgaatgc caagcagata taaatttgaa tgaggaatgt 120
 atagggtcgt gtgaagcaac ggtcgaatth tccctgggtc agtagtgaac gtgctattaa 180
 tgtaagtga ttcgtttggg cacgttcaga ttgctgtagt tgctataatt cctctagcac 240
 acaaagccc agcttgcccc tcagttgttc aaactgattt gcatccaaag cctttgtgaa 300
 aatatctgct attctgtcct caatgtcaac atgcttcagc gtgatcactt tatcatcaac 360
 aagatctctg atatagtgg 379

<210> 1124
 <211> 449
 <212> DNA
 <213> Glycine max
 <400> 1124

agcttcgtcc gcagatccct catgtaagac tatgtctaaa ctattcaaca ttatgtaaca 60
 acataattaa aacaaaaact taacccgcag atccctcatg taagactaag ttttgcct 120
 gcttcaatca agttctaagg caacagtaca tttcccaatg ctaaagtcac ctaactgtga 180
 acacaaatgg gtgatcagac caaaagcata ctaacatcaa gcattgaagg aagcattgaa 240
 cacagaatac acaatcaatt aggtattagg tatttacatc atctgttcat ttgaaatccc 300
 caactagggt gttccgccac ccattacaga agagacccta tcaataatta gcttactaac 360
 cctaggtatc tctgcaaaag ctgctcctct tgctacctcc agagctcctt tccctaaata 420
 ggcaatgtgg ctgctgtgga attttgtgc 449

<210> 1125
 <211> 459
 <212> DNA

<213> Glycine max

<400> 1125

cgccacccag ctcgcccagg cgagcaaggt ttcttctctc agaagcaaca gccttcttga 60
ggaatcttct ggagggccca agtgggtctg gttgctatct gcacccccat ttttactaaa 120
tacacccctt gccttttttt ggtgattctt ttttcgtaaa gttacggaaa cttacgaatt 180
tcgtaacgat acttgttttc tttccgtaat gttacggaac cttgcggatt acataatcat 240
cccttttttg acttacggaa tgttacgaaa cctcactaat tgtgcaacga tgcttccttt 300
tgatttccgg tgtgtcacgg aaccttacgg attgtgcac aatattttct tttgatttcc 360
ggcacgtcac ggaatttcac aaattgccta ctgatgggtg ccaagcacct taaaaatgac 420
caaacacaag ttgcatgcca ccaagcacag gtccctgga 459

<210> 1126

<211> 461

<212> DNA

<213> Glycine max

<400> 1126

gtcacctgca gctgcagcta tcatgcctc acacaatact ttgtatgtat attacaccag 60
agatgtttgt ttcaccccag gtaaaacaat atggagtat caagcatttg cccggaattc 120
aatgatggg tcaggacgag gctgaatgat tattccatgt cctatggcca attgaacagt 180
ccctttgaca agtgtttcgc acaagggtt caaggattta tattctttca ttttcccaa 240
gatacagagt gtcctatatg agagagacag gaattgtatg gtttaaggac agaccatcac 300
ccatgcaaca gtccaccagt catacataga cactattagt atatttctct cttacatcat 360
aacactgccc tattcttaag agataaaaaa gggaaaggat ccagaatcag gacgcaatac 420
ttcacatttg ttcataccaa ttcacattgg cctctgtctca a 461

<210> 1127

<211> 449

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1127

tatctgggtca aggtgaaata aaaatggtct ttcaaactaa tagttgccag attcaatatt 60

ctatatatttta atccaagtga acacttccat aagggttggt tcagtttcta tcaagaacat 120
 acatcaaata aaggatgcac agctttttaga taaaaagttt atgaactcca acatacctgc 180
 aaaagacaac tacattcgac tcgttttttt gatacactgc cacaggtctt tataacattc 240
 ctgtagaaca gcttctcaac aacaatgaca ttcagatggt tgaaaatatc aaaaccagac 300
 tgcttgagtt tatcatattt atcaatatga aatttatgaa tgtagcgctg ggcatatggc 360
 agagtccagt ttaccaatga ctntttcaaa ctacaatcag caagaccata atatattggt 420
 tcccgagtta ccacctgcac aaagtcata 449

<210> 1128
 <211> 457
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1128

ctcatcttta tttcgttcat tctcgatttc ttttcttttg tctttaacgc gctttttaccg 60
 tttatttaag ccgttttctc acctaatata tgataaaatg aatttcaacc gatcatttgt 120
 gttgtaatct cattcaatca cttttaaaac gaaatctaac cgatcgttca cgctataacc 180
 tcggttaaac cgaaaaaagt aaaataatca aaatatcttg aaaaataata ataaaataat 240
 caaaatatct tcgaataaaa taatcaaaaa aatcaatcgg acgttnttct ttggaagttt 300
 ccttgaatga attgattaat aactaaagtt aaattaagac taaaatcaac tcacaaatca 360
 agttttgtcc gaaaaatcac taaaaaccgt tttaaggtcc aacgccttaa gcggtcctct 420
 ttgcttttat cggttaacat ggaccgttca aaagcat 457

<210> 1129
 <211> 338
 <212> DNA
 <213> Glycine max
 <400> 1129

atgacaatgc ttaccaagtg gagctggccg gtgagtataa tggttaattcc aactacaatg 60
 tctctgattc atctctttgt gatgcatatg gagaatccga tatgatgact aatacttctc 120
 atgacggaga gaatgatgat gacgtgacca caagcaacgg cagggatcca cttaacgact 180

tgtatgacct atgacaaggg ctacagcaag gaaagcctag gacgctcttc tacaattgct 240
gtccatacta tgccaatata agctcaagtt tgaaagagaa aagtccatgt tgtgacttga 300
tcatggccca tatggaagac taaataaacac cactttat 338

<210> 1130
<211> 414
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1130

ttctacaatc tcgagccttc anataaccct cctcaccttc ttccgcatca ttatcatcag 60
tatgaggacg aatattacat tcttctctat gccaaatttc agattgatct atccggtttt 120
cctcattggg ctccacctca atgctatcct tttctgagct agcatggacc accttatctt 180
ctgcacgtgc ctttgtctta taaaccattt cagactcatt aaaaataaca tcatgactta 240
taatgcatct ttttgtctct ggctctaaac accacaatct gtaccctcta aaaccctgag 300
gatatcctat aaacatacac ttgatagctc taggttccaa tgtgtcttgc cttatgtgag 360
cataagcaac acatccaaac accctacgtc tatcattatt tggaggatgc cctg 414

<210> 1131
<211> 435
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1131

ctcatctaca ctgtacttct atataatttg tatgacatag tgcattgatgt ataattgatgc 60
ctatagtctc gagatgagggc atatacaaat tgctatcttg aaaacctaag accatgcccc 120
ttaatgtgat aaggggacat aacaatcaac atgcaaagga taaatagtat acatgacaaa 180
cctgtactcc aaatgtcgga cactattact tggcttctac aaaagccaag tncaattagg 240
atgtacccaaa aaataaagac ctaaattaat caacagtgaac acatcaccta ctaataaaga 300
atacaccgac aatatggtag ggaaccacac aaataaagtt ggcattttaa gtcaccaaag 360
agatgaagag aaaacatata ttgccatcat ctgctccatc atatggtaaa cacactaaag 420
ttaggaaagg aatga 435

<210> 1132
 <211> 438
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1132

agctttcttg cttcttagat ntcattcctg ttcttggtta ctgttatcgt ccattcctaa 60
 aactgccaca tcacaactta gatcttaatt gaatTTTTga gcttaccgaa ccatattaaa 120
 taatttcaga ttagggacta tactgattat ttcaactggt gttttcatat ggtgtttatt 180
 tgtgcatggt aatctccctc atgacattct gttattcatc attgtatcaa cgtaacaaag 240
 aaaacagtat tattgtgttg tcgaccaatg ccacaattaa agagggccac cctgtactat 300
 gatatatagt tacttatggt gtctttcttt cacaacaaat atgttggttg attattttat 360
 tatctatgta gttgtttggt tattcatatt tcttcctggg attgatttct tcagctggag 420
 acgagttctt cttacata 438

<210> 1133
 <211> 380
 <212> DNA
 <213> Glycine max

<400> 1133

aaccctacat ggagtgattt gctacatcta aatagtaata acatagacgt ctgttatgcc 60
 gctttaaaca cttaaataatt ataaacatac tcaaagagaa tgctgctcct tatctataat 120
 taagggttca atatgctcct tctacctgct ttttaattcac tctgattcat tcaagtgtgc 180
 ctactatacc acatgactat attagatacc tgtctgacag cttttctact tccactacct 240
 aactatatta attatgtgtc atgcagcatg actcatatat tatcacgtgt cattaagatg 300
 acatagttag tatgcatggc aatattgact tgatctaata cggatcacag tgaatctttc 360
 aaaacttaag accactttat 380

<210> 1134
 <211> 380
 <212> DNA
 <213> Glycine max

<400> 1134

atccctactc aaaacaggaa ccattcattt ttttgctctt ccttaattac ctgcttttat 420
gtgtatatca tga 433

<210> 1137
<211> 436
<212> DNA
<213> Glycine max
<400> 1137

attctttaca atcaatctat ctactgaata acaattctaa atgtaagtcc acattcttgt 60
tctttctttg tttgacatgc acatttgctc aacttcatga aaggaaacac aaatctcatc 120
ttaagcatgc attcaattta aaacaaagtc atacaccgt tttcacaaaa agataaaagt 180
gtttcactgc catgtcatct aaaataagtt aaactgttca aaatgcttca agataagcat 240
aataattatt catatataaa actagtagta tatatagaca taaaggaaat actgtacgat 300
aaccaaaatt ataataataa taaatcaaaa agtgaaaagt gtcaccagga attaaaattc 360
ctgtgactag tcctgagtct cctgtgtttg accatcctcc tcatttgtca gctgaagaac 420
tggagtagtg ggagga 436

<210> 1138
<211> 280
<212> DNA
<213> Glycine max
<400> 1138

gtagacagt caccgcttta tgagcgcttt acaccagtag cgcttogagg ccatcaaagg 60
atggtcgttt cgacgggagc gacgcgtcca actcatggac gacgagtata ctgatttcca 120
ggaagagata gggcaccggc ggtggacatc actggttacc cccatggcca agttcgatcc 180
agaaatagtc cttgagtttt atgccaatgc ttggccaaca aaagagggcg tgcgtgacat 240
gatgtcctgg gtaagggggtc agtggatccc gtttgatgcc 280

<210> 1139
<211> 354
<212> DNA
<213> Glycine max
<223> unsure at all n locations

<400> 1139

tctatataag ctgaaccatt ntatcagtgg actcaagtgg tgttttcttc gcaaactaag 60
agtctatctc ttgtatccta ctgagagtga ctctgcatat ttcttgagtg ggtcaagaac 120
accttggctg tcccaaagga ctttatcaac cttagtgggt cgccctcggg gggacgagag 180
atgctttcct tccgtttatc gtctaccttg ttctttcaaa tcacaattgt agataatacc 240
tcctatgaca ataaatatct tgtggctcta actctccttt tatgcactcc actttcgtga 300
ttgatgggcc tcaaacgaac tcaaaacgat accttgcacc tcatattgga atga 354

<210> 1140

<211> 434

<212> DNA

<213> Glycine max

<400> 1140

agcttaactg ctttgtaaaa cgaaatactc gatctatata atctttgtta tcattaaata 60
tatttcaaac tagttcaatc atatgcatca aagtatgaaa gctttcaaaa aaacatgaaa 120
accttgaagt agtattctaa acaatatttt tctgagtaca atattatgaa aaataacttt 180
ctaagtgtag tagcaacata ataagaatcg ttataacata aactaaattt gtcataataa 240
caatgttttg agagatacat ttatttatgt aatgatcttc taaacaagag caaatgcata 300
ttgacattag gttctcataa tcaagtcaaa cattgaataa tgagtgttat gactaaccac 360
ttagagagct tagttgtctt agtacttgaa cctctatgtc aagaatttct ggacaccaat 420
gtagtcttga ataa 434

<210> 1141

<211> 425

<212> DNA

<213> Glycine max

<400> 1141

atactgcatt gttgactaat tgttgttggt gttatttaca tctattttca gactcccaat 60
ttgcagattg agttttgggg ctgctttctt gctgaacaat gtttgttggga ctatgattgt 120
ataatatact tgtcttctat cgcggctgcg tctgttatat ttcttttcaa acttattctg 180
gctctgaatg catacttaag gaaatattat gttatggata atgatataca catctcttta 240

atggaacgat attacccgtg tttcaacgca tgtgttctca tgtgatgaat tgatgattaa 300
 ctgttcgcca atataataaa attattgctc tttactctaa tatatatagg taacgactta 360
 atgttgaaga ttagagaagg tgtatccctt ctactccatc accattaact atcagctacc 420
 ttctt 425

<210> 1142
 <211> 453
 <212> DNA
 <213> Glycine max

<400> 1142

agctggcttg tggagcttct atggagggtg gatcttcgag cttcaatgag gtcctttaat 60
 ggtgattttc caccatggag atgcagcgga agacaaagga gaagagggtga gaggaggcgc 120
 catccactat ggaataagcc atggaagaag gagcttcacc actaatgata gccatggata 180
 agaagcttgg agaggatgct tcaatggagg acaagaaaga gggagagaaa gagagagggg 240
 ggagcatgaa attgaaggaa gaaaaaggga gagaagttga actatgagtt gtgtctcaca 300
 agactctcat tcatcaaagt tataacaagt gttacacatg cttctattta tagactaggt 360
 agcttccttg agaagctatc ttgagaaaac ttccttgaga agcttccttg agaaaactgt 420
 cttgagaagc tagagcttag ctacacacac cct 453

<210> 1143
 <211> 459
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1143

catgtccttg ttacatactt acatagcatt aatcacaaca tacacgcaa taaatacttg 60
 gtaacaagga attataccct gcaaacaag ctatttgaaa tggaaccacc tcccctaaac 120
 ccaagtataa gtaaataggg ttaattataa aatttgctcc cttattttct tatttcacta 180
 aatggatcct ttaagggaag actagaattg tcttttgagg taaaaggat actcttttat 240
 cataaaatta aatcttatat attacatctt aagagataat atattatatt ataaaaatat 300
 aataaattaa aaaaattaaa ttttgaatta tttttcttaa aacttacatt aatactatat 360
 tttgtaacat atatttaa atgaactntg atatagtgat aaaagaatat atttcttatc 420

tcanagatca tgattcatta aaaaataaag tgatgaaat

459

<210> 1144
<211> 333
<212> DNA
<213> Glycine max

<400> 1144

accacagagt ggtacctgta gatatgtctc gggggtcattg agaacctggg gacgtcatgt 60
gggggtgctat tgcccaaaac caagcttgac caatcccgac ccaacccggg catagtcggt 120
cagtgagaac ctgtgatgta cctaagcagg cgagctcctg caatcaacag ataaacgata 180
acaagaccac aagcatggag gcttgtgggtg gctggccagc tgtgaatttt gtgtaatatg 240
tggtatgggtg cctctggtta tcgattacca aggggtgggta atcgattaca ggcttaaaaa 300
tgaagacagg atgcatagat ggtctctggt aat 333

<210> 1145
<211> 461
<212> DNA
<213> Glycine max

<400> 1145

cggagaagat gcttcaatgg aggaaaagaa agagggagag aaagagagag gggggagcac 60
gaaattgaag gaataaaaga ggtatagaag tggaactttg aagtatgtct cacaagactc 120
tcattcatca aagttacaac aagtgttaca catgcttcta tttatagact aggtagcttc 180
cttgagaagc tttcttgaga aaacttgctt gagaagcttc tttgagaaaa ctcccttgag 240
aagctagagc ttagctacac acaccctct cataactaag ctcacctcct tgagaagctt 300
ccttaagaag attcctaaag aagctagagc ttagctacac atacctctct aatagctaag 360
ctcacctcct tgagatgaga agctagagct tagctacaca cccctataa tagctaagct 420
caccctcatg acaaacaaca tgaaaataat ataaaagaag t 461

<210> 1146
<211> 347
<212> DNA
<213> Glycine max

<400> 1146

tttaatagtc attgcaccag atctaacctc tgcacagag gctggaagcc ttctcaaaaa 60
 catgaaaacc ttgaagtagc attcctaaca atatttgtga gtgttctata ttgagaaaaa 120
 tgactttcta agtgtagtag cgacccaatg agaatcgttg taacataaac ggtactcgtc 180
 attactacaa agtgttgaga gaggcattca tttatgtaat gaagttctag gcaagagcca 240
 acgcctattg acattacgtt ctcatatca agtccaacac tgtatcatga gtgtgatgac 300
 taaccactta tagagcttag ggggcttagt acttgaacct ctatgtc 347

<210> 1147
 <211> 696
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1147

acgcgctcca cntctcgccc cgtgntgttt antgcanttc attacctccc gtcgacagta 60
 atatttgatg tgtgcntctc tctacgctca tgccgtgact actcgtcgtc tactcccgtc 120
 ntagtgcact gactcatant cgtacactct cgctcattnt atcgctatca ctacatgact 180
 tactagcgta ctcttatgag tgtctctcac attctctcta ctctcttata tctatcgat 240
 ctttctcatg cacatcttct ctcatctct cggatacgta ctnnnnccgc tctcctctca 300
 cgcagatagc acatgtgtgt gctctcgatc atacatatgg ggtagaatat ctcgtagacc 360
 ttgtgcttgt cggtagagag gtcagagttg tgcccttata caatatatga cttgtttaga 420
 acagctgccg tctatgctta aatattatta tagcacatgc tcttgctttc tcgttggtgtg 480
 gacgactaca ccaatgttgt gacatgctgt atcttgcac acatattcat gtgtactcat 540
 gctacgcacg gtcctttcac gcgcttcatg ctcatgcata cctaaaatca tcatacacgg 600
 tctctcacia tgtggagtca accatacac actattatca tacctgcttc tttagaatct 660
 attgatacct ttcttgttga cctacagat actcct 696

<210> 1148
 <211> 422
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1148

cattatttga ttgaaatata tccattatat tttaaaagtc ttactctctt tcaactgcaat 60
tactcttctt caagtaagat tatctcttat gtaaaactca ttcattccatt tgggtcaaaca 120
caccatgaca ataagtttta ttatcttgaa ttcttttaca caatttcatt tgtacaagta 180
tgcattgagt tatcagattt tagtgcttga gtatttataa aaatataatt tttgtccct 240
tatgtttcta aatctgtaat tttagtcctc aatntttta ttgacatatg ttatctctca 300
ctttttataa aaaatcataa tttcagatca cccgactaat ttcaaagtt aattgtgtac 360
tttgtatttt ttttctttta attaaactta ttagtaatta aatacataaa aaaaaatact 420
ct 422

<210> 1149
<211> 429
<212> DNA
<213> Glycine max

<400> 1149

tggctgctgt catcatgtgc aagcgtcgcg atatattacg ggatttagtc agacttccga 60
gtgaaatgtt attgtcgatt gagatagctg cgagcttcgg ttggatatgg cgagcgtctc 120
tatatatgtc gggactcaat gagacttact agtgaaatgt gattgtcgta cgcattcgct 180
gctacctatg gaacaacaat tcgagcggct gacatatgtc gggactcaga cggacttccg 240
aacgatatga tattgtcgat ataatatgct gagagcctcc gttataaaca tctagcgtat 300
cgatatatta cggcactcag tcagacgtcc tagtgagatg ttaaagtcgt tcgaagtcgg 360
tacgcgctat ggctattaat tacgagcgtc acgatataat acgggactca ttcagtcttc 420
cgagtgatg 429

<210> 1150
<211> 438
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1150

agctcgtagc ctattcgaac aacaataact nttcactcgg aagtctgatt gactcctgta 60
atatatcttg acgctcgaat agtaagaccg aagctcgtag cagattcgaa cgacaataac 120

atgtcactcg gaagtcctat tgagtcctcg aatatatcga gacgctcgaa ttttaaaacc 180
gaagctcgta gcaaattcga acgacaataa catttcactc ggaagtccta ttgagtcctg 240
taatataatcg agacgctcga attttaaaac cgaagctctg agcatattcg aacgacaata 300
acatttcact cggaagtcg attgagtcct gtaatatatc gagacgctcg aaatttataa 360
ccgaagctcg tagctaattc gaacgacaat aacatttcac tcggaagtgt gagtgagtc 420
cgtaatatat cgagacgc 438

<210> 1151
<211> 508
<212> DNA
<213> Glycine max

<400> 1151

agaacgcgca tgaacatga gaacatcgag atccgagata ctctccaggc gagctgacgc 60
gtgagctgta aagatgtcac agtgaacttt ataagcttat gtgctggcca gcatagggga 120
caatcgacta tgtatagcca acgtattatc ttatgagata aggtgtgatg acggtccttc 180
cacattatat tattccatgg tagattgatc gtctccgctc aaaggaaccg cttgttagag 240
tcaagaatat tgagcgctcg tcaacgttat attataagta gacacctata gaggaactac 300
cacgttatat tgggttaatt gtggtatagg cccctatagt tgatcgaatc ggtcattaca 360
ctcatttcct ctagttatgt ctactgaact catgctttat acttttacta tgaatggaat 420
ttaacacacc tcttatataa ccgtccaatg tataacaaca ttacgagaca tggatgtcat 480
tatagatgat atgaacctac actgctct 508

<210> 1152
<211> 380
<212> DNA
<213> Glycine max

<400> 1152

aatggatatg gttaggtgta tgttaatcaa ttagacttta tccgtatcct tgtggatgta 60
taccttgaaa actgccatgt agttgttgaa cagggttcct agtaaggtag ttccaaagac 120
accttttgaa ctgtggacaa ataggatacc tagtataagg cacctgcatg tttgggggtg 180
ccaggcagaa ataaggatgt ataatccgca agaaagataa ttggatgcaa gaacaatcag 240

tggatatttc attggttata cagaaaagtt aaaagggtat atgttttatt gttctaata 300
 tagtatgaga attgtcataa ctggaaatgc aagggttcatt ggaaatgatg aaatcagtg 360
 gagtacagtt ccacgagaag 380

<210> 1153
 <211> 412
 <212> DNA
 <213> Glycine max

<400> 1153

aactcacac atatacttag ttcaaaattc tatgattaag agcatatata tactgaaatt 60
 aaattcatac acgcacagat gaacacaaca caaagggttc tctgtcagtc ggagttgtaa 120
 ttttttaggg tgttataact atgtatagtc atttcaaaa ataccctcca cttgtaaact 180
 cttatgactt aattatccct tttatatctc aagatatcta ggatgaccaa taatcaacct 240
 taattatccc cacctaattg ctaaccttac attaatactt aagttcttct ttttaagcttg 300
 taatttgat atctagacca agatctaatt atttatactt aggtcattaa tgagttgacc 360
 attatttgac caagaaaatt ctctaaacta tctttattct atgtagaagc tt 412

<210> 1154
 <211> 408
 <212> DNA
 <213> Glycine max

<400> 1154

tataacaagta ctcttgaggt ttcttccatc agagcctcgg tattgagctc ttaatcttct 60
 ttttcttctt gtattcctta ctggattcgt gcaaacttct cattcatgga tccaaaatct 120
 cattttcatt cttacaagct tgaaacatca aggatctaag atctttgttc atctaataaa 180
 atacatgtat cttcatcaac gtaaagagag tctctccaat acttaaacc taatcttggc 240
 gtctttggaa gctaaccctc attgaatgtt gtttagatgt tcaaaatttc atagctactg 300
 catatgctgg aactgtatca tgtgttgttt ctcttgtaat cttacgcaa aaaatgagat 360
 atttgagtgc caatacttac gcgtaacctt atatctcacc tacctcat 408

<210> 1155
 <211> 392
 <212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1155

gcattacatt attactcatg cttctaacca tgtctaatat tgttngattt ctgcgttctg 60
ccacaccatt ttgatctgga gaaccaagca tcttgatttg tgcaaacactc ccatgttctt 120
gaagaaactt cgcaaataaa cctgggtgctt gtccatcctc tgtgtatcta ccataggact 180
ccccaccttt atgtgatctc acgatcttaa tatgttttcc acattgtgtc acatctgcat 240
ccttaaaaac tttaaaggca tctaaagctt cattcttaga atgaagtaag taaagacaca 300
tatatcgtga ataatcgttt ataaaggata tgaatactct ctgactagtg gcattcatgt 360
ctgggcaaag tatgtctgta tgtatgattt ct 392

<210> 1156

<211> 412

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1156

tgaatgaaac accactgtca ttgagaacaa tggatccatt tttaaaaata atacctaata 60
tgtattcaca actgttgact gcttgetcat ccagaatgcc atagtattat gaatacttgn 120
gttgatactg aaacttgtgc tttcttacag ggtaggttg tgccatatat atagatgact 180
tttcatatca gtgctgcatt ttttaaagat taaaaatata cctgctcatg ctttctgtat 240
gtgttgtaaa ctacaccaat gatgtgacat gctttacctt gcatcaaata tgcattgtga 300
atcatgctat gcatgagcct ttcacgcgct ttatgttaat gcagacaaaa atatatcata 360
cacggttttc cacaatgtgt atgttactca gaccacaata tatcatacat gc 412

<210> 1157

<211> 425

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1157

ataaaaaaaaa tctaataatg agtcaaaaata ctcaaagaaa taactttgtt atgagacatg 60
attagttttt ttttatctat aaaaattaaa gatatatatt aaagaatgta taaaattcac 120

atggattatt taatcaattg aattaaactc cttaaataa aaatataatc aattaacatt 180
 tgacactaac acttaaataa ataaagtaat aacttttttt ataaaataca aaagtagtca 240
 atctaatacat cgaataattt aatattatct ttgtaattat atataggaag aagtcaaatt 300
 acaatcatat gtttttacac caacagttat attgtaaata actctatata aaatataaat 360
 tttattgatt taattgttaa attctaatta tatattatct tgagtgtggg tctcacattc 420
 tctan 425

<210> 1158
 <211> 436
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1158

tcttggtnaa gttntatgca aacaactatg ctgacattg ttgttttcta atttgtatat 60
 tgggatttat attagttttt catattgggtg ttttcaggat tcctagttag gatgcagagt 120
 ctcaactctt aaaaggcaga aaaaagtgc gtggaagctt ttgagacagc ccacgctatg 180
 ggtgttatta tgtttgactt gccaaattgc cctaagaaac gttgtcaact agagacatcc 240
 tctgttaatg gagaaggatc atctacccat actgttactg catcttttga aactgccttg 300
 aggtgggcag ataagtagtt gatgcagtca aaactgcatt tataaggctt gcaaattgtc 360
 cttcttttag tatcggtgaa ttcgaggaac tactaagaca aattagtaat ctggacttgt 420
 ctgagttctc ttcaga 436

<210> 1159
 <211> 426
 <212> DNA
 <213> Glycine max
 <400> 1159

gcacttatgc agtaaagaga ctgtctaattg catgtttaca acatggctaa gttcgactaa 60
 aaccctcttg tacactttat gtgcaagtgc tagcagacct actggtgaga aaaacaaatt 120
 tcacacttga caatggatgc acgcgctgct atcgctatgg ccacataact tcattcgact 180
 gttgtaagca tctacagcat gtctttgcaa tataatgtta gatactaaca cttggacagc 240

tgtatctatg cgttgacggc tgagagtgcac cacattgatg gatatatcta tcacgaataa 300
 gcgtatatta gaagactcga ggccgtgctg gactacaaag tctagttgtg gacagaaaca 360
 tgctcgggtct gaatggaatg ataagaacta aaatcaatca ctgacatgag cagggatgat 420
 catcat 426

<210> 1160
 <211> 502
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1160

cgcgcttgca cattgtgacg cgtttcgccc gcgcaccttg aggcagctgc ggtctacagt 60
 catatgcttc tggcactcta cctcatatct gtaatagcca gtgtagttat ataaagcctc 120
 aagctcatga ccgtctatct atgcgcgcgc gtatgatcac ctgcgagacg tttgtctggc 180
 caccgatgat gtcattggaat gagaagtcac gcttgcccta tctgattacc aacacgcacc 240
 cctcctgttg aggtgcttgt gagcgcctatc aggtactgat ctcaaacaga gaaagaggaa 300
 ttgcaatgtt gcggatcaaa tgcttgagct tcaactgcta cctaacagct tgcactgtgt 360
 cattcatatg ttcgtccaac cgtgcctcgt aaccgaacgt aatatgagct ccatcatctg 420
 tgagtagcga gtgaacatta tctaaccncg acttatgtgc aaacgctatg ggatgactac 480
 tccaagatat cacgagagag gg 502

<210> 1161
 <211> 380
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1161

atctaattatt catgtgagtt acagaaaccc gcacttattg tgatcacact ctttgtgata 60
 cattattatc attattaatt aaaacttata aaatcgagaa ttaaattcca ctgatgaata 120
 agatgtgaag ccgacnattt tttagccttt cataaattgt aacagttcgt acaaatcaaa 180
 gcatattaaa atgtgtttta gcattttctta caatacataa tgaaagtggc ctctagtc 240
 gtacaacaat ttgtcaacta gggcaagctc acattaaata tatgggcatg gctaacaatga 300

ttatcgcaaa gtttctaaca ttatatttac ccgtaaacta ttatgaatgc ataacagaaa 360
talcaggaat attggtgttg 380

<210> 1162
<211> 424
<212> DNA
<213> Glycine max

<400> 1162

agcttttaaaa gtacccattt accaacctca tagtctactt cacgccattt agtatcatca 60
gtgcgtttca tcgtagcttg cgcttctaata agcttctgac gaatgcaaag aaaaacttcc 120
tcacagtgtc tcagagtatc atccaccgca tctagtttgg aggagctagt gatatagtct 180
ggaaaagaga aaggtttccg cccaaacgtt atctcatagc gcgtagagcc cgtgcctgcg 240
ttccatgaag tattatgcga taattcgacc cagggaagga atctgcccc agtccccgac 300
cggcgggtgca ccatagcccg caaatattgc tctattactc tgttcatgac ttcactctga 360
ccatcactct gtggatggta ggctgaactc attctaagcc ttgtaccact caattaaaaa 420
agct 424

<210> 1163
<211> 427
<212> DNA
<213> Glycine max

<400> 1163

taggtgtttt acaaaatcta tttagttaaa tatgctacga tttagtttat aataaaacct 60
attaagcttg ataaattggt ctatttattc atttatatat aataaaaaat taatatacat 120
gtattatact ttaatattha atatcttaat aagttaataa ttcatatcat aaataaaata 180
aatatttgag ataaaaagcc tttaaagtaa taataggtca talcaggttt ttaaaaaggt 240
caaaccaagc ttaaaaaaag tctctgatag gataataggt taggtcaaac cttaattttt 300
tataataggt caaacctatt tacacagagt ctaacatgcc ttgtatattc tcaccctat 360
tcttacgttt caatgtttga acttcaaaaa gaaaaacaat attcataata tttgcttcca 420
atcatgg 427

<210> 1164

<211> 463
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1164

tcctcaacat caaagtaata caacatccaa tcatcatgga ctatcaaaat caagcaaaac 60
 agggcaaagg cagaaaactc tgcccaaaac acaactcana atcacagctt ttctcactta 120
 aagacccag taaaaattcc ttcgttcag tttgttaacc gttggatcga ctcgaaaatt 180
 ttactggaag tctctagtag ataaacctac attntgaccg ttgggatcta ctagaaaata 240
 tccagaactc cttctgcaat attctttcca cagccaatca cacacaagca tttttctgca 300
 cttgtgcaaa attctgctgc acaatttcac agcaaaaatc tgcacaaaga gcagatttcg 360
 aaaaccacac ttccctcat ccaatctttc ccaaatacaga tctacaagt cccaaatcat 420
 gtatcaatca tgtctaaacc aaagtcaagc ttcaaaacac agc 463

<210> 1165
 <211> 401
 <212> DNA
 <213> Glycine max

<400> 1165

agcttggttc gaggtactca cccgttgaag atcgatgatc gactatgaac gaatgaagag 60
 cgctgaataa cgggtgaaac ctttgcgaga ttcctcacgg aatacgttac ggaaacgttt 120
 cggaagcgcc tcggcttaga ttgtcttcac ggaaacaatt tttctagca tattcgaaag 180
 agagagaagt gcctattgtg ctgaaccctt ttcttattgc cttctctccc tatttatagc 240
 taaatagggg aggtggttgc cgcccagctc gccagggcga gctcaactcg ccctggcgag 300
 cagggttgct tctccagaa gctaccgct tctggaggaa tattccagag ggccaagtg 360
 ggctgggtg ctatctgcac ccacattgta ctaagtacac c 401

<210> 1166
 <211> 460
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1166

cgctaagcga gatctgcatg atttaaatac attcttcagg gtcaaaaaca caattttctc 60
atccccctct ctcaaaaactc caccaaacca ccctagaaac tctctctcca ccaccagga 120
ccatcggtag ccaccacaag ccaactgttgc ttgccgcaa accaccatac ggagaggaaa 180
ttaaagtca cagcggacat taaaagagta gggaagaatg agacaaacac acaagagtnt 240
gtatactggt tcggtataca cccgtgccta catccagtcc ccaagcaacc tgtggctctt 300
gagatttctt tcaac'cttgt aaaaatcctt ttacaagcaa agatccacaa gggatgtacc 360
ctcccttggt ctctttgaac ctagtggatg taccctccac tagaactgat ccacaagaga 420
tgtaccctct cttgttctca gtcaataacc caagtagatg 460

<210> 1167
<211> 586
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1167

gcaaaactcc caccatacgc ttctactaat catacatcac tctcaatatn acatanaccc 60
tcatacacac gccgtgcgtg aaaccattgt aganccgtcg tatctccaaa gacgacctcg 120
angcatgcaa gctgctactt atgcggcagg gcggacttac tttactatct tgtctccaac 180
gcgagctatg accactgttc ttccttcccg cgacgcttct tttcatgtcc gcctgagtgg 240
gtatatagcc taaaccatac tttccacgat ctctttgagt attcactcag gctagtattg 300
ctgacgctgt ctttgccctaa acccatcccg ggtgcataac cgctacacca acataactag 360
ggccattatt accgctgcat cggacagact aggctgctca aagagggagt ccacggagga 420
tatgtcgacc acctacaaag actgtagagc gtgctctaac gaatctattg cggatcacac 480
aaatgcaagg tagacggtca gcttaacaat atatgataca tgcccgatac gaagcaaaca 540
atgccctccc actaccaaatt attagcatat gccagacgc aatccc 586

<210> 1168
<211> 380
<212> DNA
<213> Glycine max

<400> 1168

taaacagaca attttaaaga catgaatgta atgagatttg agtcttgatt tgcctaattg 60

cattcaaaac tctcattgga aaagaaatcc atgtctatga acttaggata aataatggaa 120
 tgagaggaga aaaggggttg gtaccatata cgttggttct ctgatgagaa catcaaggaa 180
 gaagatatgg acgacggaat ctgcatttcc tgagcctcgg agtgccggtt ggcttcactc 240
 gaagatccct tgcacttttt tgatggatct gccatttgaa cgagttattt gaaatatcaa 300
 tcgggtcacg tgaaagagaa tgacaacaga tgaagtttgg gctttcgggt gagtgatttg 360
 gacaacactc tactgatata 380

<210> 1169
 <211> 427
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1169

tgtttgtcat catcttaaag ggggagaatg tgaatgtatg tatacatgat tctgatgatg 60
 tcaaagaaga atctaacaag gctgcttcaa atgataagca tttgcttcaa gaataattca 120
 agattgcttc aacaaacaaa gccttggttc aagattcact aaagaccaag ccttgccctta 180
 aaacaaagtg ctttcaagac atgcaaggct ctggtaatcg attaccatga agtgtaatcg 240
 attaccagaa gacaggggtg agaaatagct gttgaaaaag gttttgaatt tgaattttca 300
 acatgtaatc gattaccata tgtctgtaat cgattaccag caacgaaact ttggaaattc 360
 aaattcaaaa gtcataacct ttcaaattat aactgtgtaa tctgatacac aaacattgta 420
 atcgatn 427

<210> 1170
 <211> 462
 <212> DNA
 <213> Glycine max
 <400> 1170

agctcggttat taaatacaaa acacacatat tattatgaaa aaattgacgt taatgacgta 60
 aattattatt aacacttacc actgcatgtc tcagctagtc gacatcagac cttgcagatg 120
 tcgacgggtg tgctgcctcc gtgaccggac ggatatctgt gtctggatcc tgaggggcaa 180
 ctctgggctg cgtagcatga ccatctgccc gaggatctga tggctggccc ggcgtcatga 240

atgggatgcga catgcggaag aatcagtcga tgtaatcgct ggcacactgc cctggcacia 300
 cgcagatgtc acctgctaca accatatggt ccgaatagtg catccacctg ttgtgtatat 360
 catcagacgt gacccatgaa tcggcaggtg gagcatgaat ggtctgagtg tatcaaactg 420
 ccgcatgacc ctctttgggtc ggtaatatata aacaatgggc cc 462

<210> 1171
 <211> 512
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1171

cgcgcgtaa tccgttggtg annaccgtgc tcgtaccctg ggatactcta gagagatacc 60
 tgcaagcacg caagcatatg atactgagag atggaaaaca ctattttaat gacactaata 120
 gggacatagt gttgtattgt attacaccac gtaatcccca tgagcctagc acactttctt 180
 aacttatgct ctgcggaatg aactataccg ttattaggtg agccttgaac agataaccac 240
 tccctctagg aacaactttt tgctcatgat gcactatggc ctgtgtatag aagctatgtc 300
 tcattcataa aatatgcttg aatcttgtat gctacgctac acctcactta cacacaatac 360
 ttttgtaata ctttcattaa acgccaatcg tttcatttgc cggtagattt attctgctta 420
 gcaccctaca tcccttaca aaataggcac catttacttt tttgatccta cttcatcacc 480
 tcgctctatg acggctatca tgaacaataa cg 512

<210> 1172
 <211> 390
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1172

tgtcttctgt ttatatatga tttatactcg atctaagact tgtctgatgc aatttgctca 60
 atcntggatg acggcaatgg tgatttcgaa aatctgcact tatatgcaga attttgctgc 120
 ttaagtgtgc atcgtaatct tgtgtttgtg cagaaaatgc ttatgcatgg ctgtgtgtgg 180
 aaagggttgt acatattggg gtctggacgt tncctaacat atcccaacgg tccaatgta 240
 gacttatgct ctatggacct ccaactcaaat tttcgagtct atcacacgat gaacgagttg 300

gaacgatcag aatgttactg gggctctccga gtatgaaccg ctgcgggacg tgtttgtgtt 360
 ttgggcaggg gtctttgcct ctgccctatc 390

<210> 1173
 <211> 428
 <212> DNA
 <213> Glycine max

<400> 1173

gggaaataat aatctgccgc agcggatcca agagaaaatt aagagaagaa ggagaccctc 60
 tgaaggctat ctatgctacg gtatgattaa cctttaccca tcgactttac cttttatgcc 120
 aattgtgaat ctttaatcat gaaggttaag agaaggagcc tgtgtagtgt gtagaggagg 180
 gaaaataagg gatgcctata aaaattatga atggaagctt ataaaatatg aagtacaaat 240
 acatacaatg ttgcacagat ctaaagataa tggcgaagag tgtcattagt tgatactttt 300
 ttgttatact atttctgtat tctgtttgac ttttatacgt ttctgtcagg cgtcataaaa 360
 aagaccgcac atagctagga agactggtgg atgcgatagc ctatatagca gtgataggca 420
 taaacaca 428

<210> 1174
 <211> 118
 <212> DNA
 <213> Glycine max

<400> 1174

ctataccac cgagcctga gatacgagc catttgcaaa catcatatac tacttgctgc 60
 acaactatac ttgcttatga ttgccaaacta gttattacat taatgtaagc tttctgtc 118

<210> 1175
 <211> 483
 <212> DNA
 <213> Glycine max

<400> 1175

cgcgatgaac catggatacc gtgaataccg gaacctcgga gactcctgcg gcgtcgagtt 60
 aaaaaagacg agtttgcatt tttgacgggc gaacgcgtcg catgatcatg aaacctagac 120
 ctccctgcac taatgtataa ggatgctaaa aagcatacta ttcgttccaa caatgactct 180

aactgctgt atctcacctc gccttggtgc atgatttact gactaccacg cctgccttt 240
 gaaaggatgg atttcaacac ttgtcacgct ctggtaatcc gttaccagga agagttatcg 300
 atcaccagaa caccagggcc ttaattatca gctcaatcag ggtttgtatt tgtatctgca 360
 acatgtcatc tgataacacc tgactgtaat ttagtaccac tctgagctt aggaagctac 420
 gttctaacaa caaaactcct caaattattc ggggggtccg acacaccatc tctgataccg 480
 gcg 483

<210> 1176
 <211> 424
 <212> DNA
 <213> Glycine max

<400> 1176

ccaggcgacc tatgttgctt cccctagaac gcactgtctt ctggaggaac ttcttgaag 60
 gcccaagtgg gcctgggtgc tatttgcacc cctgtttac taaatacacc cctgccttt 120
 ttttgctgat tctttttccg taacgttatg gaactttacg aatttcataa cgatacttgt 180
 tttctttccg taatgtcacg aaacettaca gattacgtaa tcatcccttt ttggttcc 240
 gggatgatac gaaacttcac ggattgtgca acaatgcttt cttttgactt acggcatgtc 300
 acggaacttc acggattgcc taacgatggg tgccaagtac ctcgaagtgg tctaacgagg 360
 gtcgtcatcc aacaaatata tgggtccccg acgatatatg ggtatgacag ttgcccctct 420
 ttat 424

<210> 1177
 <211> 401
 <212> DNA
 <213> Glycine max

<400> 1177

tctacttatg tggcacggcg ggcttacttc actttattgt cttcaacgcg agctctgacc 60
 actgttcttc cttgccgcga tgctactttt catgtgcggc tgagtgggct tatatcctaa 120
 accatactta ccacggtttg cttgagcatg tatcaagcta attatgtccg cgctgtgtat 180
 gtctacaccc atatcgggtc ataaccgcgc cccaacatag ctcggggccat cattaccgtt 240
 gaatccgaca gacaaggctg cccatagagg gagtccacgg aagatatgct gaccacctca 300

taagactgga atgctgttct aacaattata tttgagcttc cacataacgc atggatgatg 360
ggaagcttac caagatgtct tctctcctg acacgatgac c 401

<210> 1178
<211> 432
<212> DNA
<213> Glycine max

<400> 1178

agcttctgaa gaaggacttt actactctgg cgcagactag gagcgtcttg tctacttca 60
accttgccct tacatcacat acatcttata tgaacttaga tagggcgagg ttggtgtatg 120
gactagatat gaagatggat atgaatcttg gagccctcat ttctggacag atatctctga 180
tagctcagtc caactcctcc cggctaggat ttctagcgat tatcactgct ttatgcatgg 240
ccacaggagt caccttagac tcgttgactt tcaaaactct cagcccagct attaacttgg 300
cttacatcaa gaagaacttc tagaacttgg atgacccttc ggtcagcttc ccagggaccc 360
gtaaggccag ggccagagga tctgagggtc catcttcaac tgctccccag gactctacat 420
gtccagctcc cc 432

<210> 1179
<211> 427
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1179

tagtagcagt tacagaagtc aatatttccg ccaataacta tgatcatttc taccttacct 60
attttccatc atggccttgg tttgtgtgtt tagattgcca ccagagtcct aaatagacaa 120
tatctttcat tgcaagctta gcaacagtc caaaacccaa tttttgccga aaccaagtgt 180
catgatttct atattaccaa ttttgctagc tggtgatgtt gcatcatagt tttgctatgt 240
catctacctt tgggtctcatc tctttacctt acaattcagg caattatata attacccttt 300
ttcaatatat agaattggca acatgcaaac atatctaata caggaaattc caccactaat 360
agtcagccta taatccataa ccaatgaagt ccccatctc caatttattt catcatctaa 420
ttntatt 427

<210> 1180
 <211> 407
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1180

ntcacttcat aagttgtctt atgcctaana atacgttgag ttatccttag taaagggtaa 60
 aatattgact caagtaaagt tcaattccca ctcaagattg gagcaaacta cttagcttgg 120
 caataattcc gaccaagggtc caacctagta ggtaaccatt cttgactgtg acgtatcaag 180
 gaatgtattc ttaacgaata aagttacacg gttaagaatg ctgccgacag caacaacaaa 240
 atggaaaatt ttgtttgtga caggaaaatc ttttgtggac caccatattg ttatttcaat 300
 taatcttctt tttcatcgtg aaatcttttt ctgtgctgct ttcccatttg actctggaaa 360
 atgaagaagc tatatagatt tagatgcagt tgtttattta tttattt 407

<210> 1181
 <211> 345
 <212> DNA
 <213> Glycine max

<400> 1181

gcttcaacgg cgatttacca ccatggagat ggcgtggaag acaagggaga taagggtgaaa 60
 cgaggcgccc tctactacgg aataggccat ggaagacaga gcttcaccac caagaatgtg 120
 ccttgataa gacagcttgg agaggatgct tccctggagg aaaagaggca cagattgaca 180
 gagagagaga gagagaagat cgaccttgaa ggaggaacac ggggagagaa gttaaactct 240
 gagttgtgtc tcacatgact ctcatcacc acatttacga caagcgatac atgtgctgct 300
 atttatagac tacgcagcat acttgagaag cattcttgag aaaac 345

<210> 1182
 <211> 421
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1182

tccatcgagt ggtaatcaga gcacaagagc ttotagcaag tgctccttac acctccatta 60
 attttttttg ctttaccttc tttccattg ttgtttcttc atttttctcc atgtatctcc 120

tgcataattg ttatactaac tatattcaat gtacctgaac aaaatgattt ccaaacacgt 240
 gaccgacaca tatgatgcgg tggccataag aatcaggtgg tgtgtgactt ctataaggga 300
 aaaatgtcat gtcttggttg cgggacaacg atacaaggat tacgttatac cggaagcaa 360
 tcacatatcc catgtccgtt atattcatcc actcgtccac gcttacctga atgaaccaa 420
 catacacatg taagctaata taacatt 447

<210> 1185
 <211> 388
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 1185

agcttcttat caaggctcat cttggtggtg aagctccttc ttccatggct tattccttag 60
 tggatggcgc ctctctcac ctattctect ttgtcttcg ctgcctctcc atgggtggaaa 120
 atcaccattg aaggacctca ttgaagctca aagatccagc ctccgtagaa gccccacaag 180
 caagcttcca tcaagtggta atcagagcac aagagcttca agtaggtgct ccttaaacct 240
 ccatntaatt tttgctttac cttctcttcc attgttgntt cttcattggt tctccatgta 300
 tctctcaca tgtcttggtg taaatgtttt aacatgattc tatagagctc ccaactgatta 360
 aacttgctat acacgctaga ttgattt 388

<210> 1186
 <211> 347
 <212> DNA
 <213> Glycine max

 <223> unsure at all n locations
 <400> 1186

catagatgga tgaatgtatg ggacgaggtg gtacgctagc tctagttcca taatggagac 60
 cttctgctga aactaacttt ccacactttt gcacttcaca cgaaagagcc ccgcacgaaa 120
 ctaggataga catgtccatg taaagacaga cgctttccaa ggaactacct tccgctcccc 180
 atcatatatg gcaacgttct agattgctgt gcaccctctg atcttctagg ccattactgg 240
 actgataatt ttgcgcgagc gacgacttca gctcatggac taccattata ctgattttca 300
 tgaggaaata cggcgcctgc tgtgggcatg actgtttact cccatgg 347

<210> 1187
 <211> 376
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1187

taagtcacct gcggcatgca agcttctaaa ctttatacaa gaatgaagct ctgataccac 60
 ttcttggaca agttgcctca gatattctta gaaggggggg ttgaattaag atatcacaga 120
 ctattcccca attaaaaatt ctacttttta tttaatccaa caaccaaga ttccttttaa 180
 acaagaactc ctagataata atgcaaatta atcttactaa atagaaataa taagcaataa 240
 acaataaagg agtctaaggg aagagaaaat gcaaactcag atntatactg gttcggccac 300
 acccttgtgc ctacgtccag tccccaaagc acccgctaga gaggttcact atcttgcaaa 360
 atccctttac aagttc 376

<210> 1188
 <211> 415
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1188

gacactctat agtactacag ctnttgttct gttacatctg tttcatgtat atgatgtgat 60
 ctccaaatat ccaccgacta gtcattctta accagtgcaa ttttacctga tatcttggtc 120
 ctgtagaatc tcatgtcctg ccgaagatgc ctgttcaaaa ttgcccgtta cagtgaagaa 180
 tcctaggaaa aatataagaa aacttgagga aaaagtgata ttatactctt catttaaaag 240
 tagtacaacac tacaacttta agtttttagct tttagagcat attcattctc caataagtga 300
 aaataatttg gtgcagcatt taggaagaat ttaattttct ttgatcggtt tgcaggatat 360
 tgctggattt cataacagta tttaaattat gcaacagatc aaacatgggc atatc 415

<210> 1189
 <211> 412
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations

<400> 1189

agctnntgcg gattggtctn tgccagtgaaggatcgatg tgggtccgaa aagaggcaaa 60
tntagtcatc ctgcttcgac gaatgagaaa actggngcaa atgaagaggg tgaggatgaa 120
agagaaatcc atgctgcatg taccattcct acatggaaac ttcccaccaa cccaacaatg 180
tcattactca gccataaca acccatctcc ttaccaccaa cccagttatc caciaaggcc 240
atccctaaat caaaccacaa aaccaccta ccacacaacc aatgctaaac accacttttg 300
gcatgaaccg aagcaccaac caaaagggat ttttgagca taaagcctgt aggatccacc 360
ccaaattccg gtgtcatatg ctgaacttgc tctcatatct actcgataat tc 412

<210> 1190

<211> 470

<212> DNA

<213> Glycine max

<400> 1190

tcagccagat cgctaagtga gagcttatcc gtggctaagc atgacctatt gtcgccaagc 60
gcaattcctt acgaccataa ttgaggcca tgacgctaag caccagtcac ggcagctatg 120
cgagattcat tgtggcaata tgagcgctaa gcgagtcct ctcagctaag cgcatactcc 180
tctgtactta agatgcatca ttttagctaa gctggccaga gcctgtttta gcgagagttg 240
tagcttttct aatctacaga cctcgctaag cggacatacc ctcgtgctaa gtcgagtttc 300
tgctaaaaaa aaaactgatt ttgaatgtga aacgtcagct aagcgcacgg gtccgctaag 360
cgagccttgt tgagaaacca aacgtctctc ttgctcgctt agcacaacgg tccgctaggc 420
gaaagtatcg aaaaactgtc taagtgagtg taacagcagc tacactcaca 470

<210> 1191

<211> 474

<212> DNA

<213> Glycine max

<400> 1191

cggttggtgca tctactcgac cggatcttaa gtcactgggc tgcagctcta gccaatggac 60
taccttgatt aattcctttg gagccctttt gagccgtggg tgcccttccct tggtttgaag 120
ctcactacaa gccctaaatg aaaaaccatg atatcaccct atctttgacg aattttggag 180

[illegible]

<223> unsure at all n locations
<400> 1192

<210>	1193
<211>	504
<212>	DNA
<213>	Glycine max
<400>	1193

493

ctcacatgtc ttgagactaa tgtcctaaac atgatattta gagcgaccac tgattataat 420
 tgctatgtaa gcataatgtg atcgttatag cgcacatctc ttttcttgta tctaacaatga 480
 attggtagag taaaggcctt tagc 504

<210> 1194
 <211> 417
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1194

agctntatgg ccatgtaaac actatggcctt atggtttggtt ttccccatt caatcaaccc 60
 aatgtttcca aaaaacatct cttttatcaa gtcacgcaca catccgagtc catttaggca 120
 tccgggaaaa atctttcatt gcattcacc ttcaggcgca cacatttttt tcaaaaatct 180
 ttttatatcc taatctgtga attttccaaa gaaaactggc ggtcattttc tttcaaaagc 240
 atgttggcctt tttagttttc tttctcttag ctnttttttt caattaattt ctttcagacc 300
 aatttttttc agaaaagggtt tgtaacctgg gcaaagttgg tattcgagat tacactntat 360
 caaaaggaac aanaggcgtg tgaatgacaa taaaccaaca cacaagacc ctcttat 417

<210> 1195
 <211> 426
 <212> DNA
 <213> Glycine max
 <400> 1195

tcttcgaagg gcaagggttat ttccagtttc ttgaaaatat ctaaaaatct tgccagatga 60
 cgatcttttt ccttcttgga aggtaccaca ggatatagta cttccacacc ttcacccaca 120
 gctttttcac ttctactctt ctttgcattc ttattttatt tattcttttt cattttctat 180
 ttcttattct acttcttttt ctttttcttg gtccttcaat tctttattct ggaccattat 240
 ttgtttccct ttttcccgat tgccttcacc tctcacatca tttttcttaa cttcagtacc 300
 tttcttttta gtcgctttct ccttgtgcac tacactttct tcacctcag cttccacaaa 360
 cctcttactc cttgccatca cagctttgca ttctctcttg ggattctgtt ctgtatttgc 420
 cacaaa 426

<210> 1196
 <211> 464
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1196

tctcccgcat tttcctataa ttagggggag aagtgcacagg aangaacgtt caaccctcct 60
 ggtatatgag attcacttac tattagttag ataaatcgct tccatgaaga atatacacgc 120
 cgaggcgctt ccgtaacgtt gatgttacgt tttcgtgagt gatttcgcga agattttcaa 180
 ccattcttcg acgttcttct tttgatattc gtcgttcttc ggtcttcaac cggtaagttc 240
 ccgatatcga actttntaat tcattctatg taccgttggt ggtccccatt cgttttagcgt 300
 acttttattt tcgtttcata tactctacgt agctcctttt gacgtgcttt agtcatctac 360
 ttgcctaatac aataataaaa taaatttcca ccgatcattt gaatgggttac atcacttaat 420
 ttcagttcaa tgagatgtga ccgtttgggc atgccataac catg 464

<210> 1197
 <211> 100
 <212> DNA
 <213> Glycine max

<223> unsure at all n locations
 <400> 1197

acatgttgct cccctatct ctaacaagct tctatagttc ctgtgatcta tatggatact 60
 gngatagcta cttatategc cttgcgttga ggacgggcta 100

<210> 1198
 <211> 281
 <212> DNA
 <213> Glycine max

<400> 1198

ttaacatcct cattgattag tgcagatcaa actactatat ccttgagggg acttacatca 60
 tacacttgag ctgcattttc cataaaccag aattcgcaaa ggatggaatc aaggaatagc 120
 atacataaag atcatgagca taccacggtt ttcaaatttg accataaagg gccataaaaa 180
 taattcttat atttctttta aagatatatt attattaaat aacacatagt tggatatagaa 240

tggtgatgct tacaagagtt aaaaaattat tctatccaat t

281

<210> 1199
<211> 197
<212> DNA
<213> Glycine max

<400> 1199

taagtaaacg atcataaacc cataatctgg cgacaagtgc agataatgag agtcatggct 60
agttggcata acatgttaac caatgcatct agttttacctt cacgcttcct atttctgttg 120
atgaatatga attcacggct acttgattca ctctctaat gaccatagca tcacttctgg 180
cactaaattg ttgggag 197

<210> 1200
<211> 413
<212> DNA
<213> Glycine max

<400> 1200

cacggagact aatcagacat gggatgcagc tatcacgtac atgcttctat tctaaaactt 60
ctcatcatgt gcttattagc tggtcggggtt tctctttggc tattgaagcc ataccaatta 120
tggacaatat tatggtaaca ggcagaagat ttgcgccgatg attcattctt gggatactat 180
ggtaagaact tggaatctca tcttcaatga gtcattattgt cataccatga aatatcatat 240
cttggttgcta ttgaatgaat cttgtaatta ctacaaaaca cctactgaaa ttctaaattt 300
tttagatcaa atgctaagta caagtaaaaa gatggatgat gatactattc tgcaggctctt 360
agctgtgctg tgtacgcggtt tcctataatt gctattgcta tcattggact tct 413

<210> 1201
<211> 596
<212> DNA
<213> Glycine max

<223> unsure at all n locations
<400> 1201

cttcacactc ttntcaacat ttaantgtgt tgtntntant taatttannt ttttnaggcc 60
ggaatgatgc ctcatcactg acannctaaa gtaaaccgca gcattgagact taacaaccag 120
tagatatgag tgaattaaac tacgattaca agaccttgaa caaatgtgga agaatatgat 180

tgaagcatac tgttatattc tagcatatct acacagtaca ataacgtaca cttaagaatt 240
aagccttagc ctttcatctt tggctgctgc attattcgtg taaacatagc gcatatgcca 300
tatactaacg ccgtattact gtaagtgaac tcggaggcat tacaagaaat gagaattgag 360
acagttctta gtttcgagtt gataagataa gtgcaccatt acaaggtgcc aacatttggt 420
cacaccaac cgaaaacaat tacagcaata aaaacaaagg tcatgtacac atggtatgtg 480
attatagctt tgtacatgga tggttctaag atcaaaaagt acgcttgcca tttttacgta 540
atgtggtatc aactttggtc ttaggccgac cattaccaa cactacaatt gtcatt 596

<210> 1202
<211> 484
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 1202

ntngatgctc cgccagcaac tccgtaaggt actgatcgaa acctctcagg atagattagg 60
catcttttat ttattttaaa cgaacaatag taataattac tgtgaattta aaggatactg 120
ggctgcttta tagcagcgcc ttcgcttgt cccgggctag gcaaaggacg acgacctact 180
ggtcgtgacc ctagcctctc tttgcatccg tccataagta cctaaaagta ggaaacaatg 240
aggtgtggca aatcgcgacc ggcgtcgtcgt cttaccttaa tcggtttctg cttttaactt 300
tgtctcaacc tttgatgatt ctgccccctg tttatcacia aatatgcatg tgtatgcgta 360
tgcataaatg ttttcaaacg caacaaattt ttagtgaaag ctggtttagg ttcgattnta 420
attaagcgct tggggcatcc catgaactga gcgaaagggc tcaggtgatc acaaactaac 480
acat 484

<210> 1203
<211> 441
<212> DNA
<213> Glycine max
<223> unsure at all n locations
<400> 1203

ccaacgctct gatcaagctc tcccataatc tanangtaaa tctacgatct ctatcagata 60
ctatgctaga ttgcacacca tgtaacctga caacctcact tatatacgag gaggtcaact 120

tttccatgga aaatctgata ttaatgggaa tgaattgagc aaacttattc aatctatcaa 180
 caataaccca tatagaatct aaacctctaa gggttctatg tagtcctacc acaaaattca 240
 tggaaatgct gtcccacttc cactatggta tctctaaagg ttgcaactta cctgaaggtc 300
 tctgatgtta tatcttatcc ttctgacaga ctangcatgc atacacaaac tactaacct 360
 ctctctttat gtatagccac caaaacatcg tctttaaatc atgatacatc tgtggagcac 420
 catcatcaat gctcaaatta c 441

<210> 1204
 <211> 313
 <212> DNA
 <213> Glycine max

<400> 1204

taatggttgg atttgacatg ggatcactaa ccgcacatgt gtggctcata gtgccgtcca 60
 catgctcttt acaatgtatt gagaatgtta ccgcctgccc ttactagctt ctgatgcgca 120
 caagagagct cgaggatcat ctgcttgtat acaaggggta tgctgtaatg catatcagat 180
 agtctgttga gcgcgtatgc acgacgatgc catgactgct gcgacactag atgctggtgt 240
 ttgataacag acatgagcac gaatgatagg ataaacgtga tgtgattaac agatgcttat 300
 gcactgcatg ata 313

<210> 1205
 <211> 239
 <212> DNA
 <213> Glycine max

<400> 1205

agttaactga ggcgtgcgcg cgcgtgctct gaactagagt tcaacttact gctgagcttg 60
 cgcagatcat ggaatatgca gatctaagca gaagctcgaa gtgattcatg atacagacca 120
 tagctctact aagacgaaat aaataaatga gtatcaatca ctccaatgag catgctctat 180
 ggtgacatca tggattatca cttggcgtga tgaagtgate tgctcgctta tatattaat 239

<210> 1206
 <211> 357
 <212> DNA
 <213> Glycine max

<400> 1206

attctataga gagaaaggtc cacgttccac agagtattga gagattctgc agcgtgaaga 60
tctgtacaga ctagagtcgg aagcggaagc cgttctgaga gctggagatg aatccgtgac 120
tggttgtgag atcctacagg tgaaggagac atcctcagca cttgtatttc tgcgatctat 180
cgtcttgttc atagctttcg tgtaaaggac gctccttgga tatggaaagc tcgaatacta 240
tgatggatct tcgttgtttg tgcacgatgt ctatataaat ctatgtattt aatgaagtta 300
cgtgagatgc ctgtgctata aacattttta ttcagtatgc ctataccatg ttcatat 357

<210> 1207

<211> 458

<212> DNA

<213> Glycine max

<400> 1207

tgctgcgaaa gagagttttg ctgcttattg catgcctata tagctctaca acataattaa 60
gagttgtgct gaaagatatg tactccttct gcctatgtaa attatgtttt atgtgattat 120
ttataactta tattgggaat tgcgatatat tgcagagtt ataaggctga tattattggt 180
atcattatth tttttttttt gaaaattatc tctttttcta tctcgtgttt aggagaattt 240
cttatactat ttacttctct ttagtagtac ctattgcttc ctttgttttt ttttttttac 300
ctcttgaact cctccgattt cgggtctctg aagataatat catcgcatgc tgttctaaca 360
atatacatgt ctgtcttata gtcattattc ctacggagat gtatggacta caatataatg 420
gctaagcgta agaagaggta aacactatta tatataat 458

<210> 1208

<211> 517

<212> DNA

<213> Glycine max

<223> unsure at all n locations

<400> 1208

cgggttgatt ggatcctact cgtacccgcg atcctctgag tcacctgcgg catgcaagcg 60
ttgatagtaa aaggctgaga tgaatgtgca accttgatat gcagaccatg atgaagagag 120
cctntgaggt accttacaat gcgtataaca gctgtccaat gagagtccat atgattagcc 180

atgatctgac agaccttggt cacagcatag cttaactcat gccttgtaat ggctgtgtat 240
 tggagagcac caacaacata cctatagaga gatggatcac tgaataaatc caagccagct 300
 ttggttaact tgcaattagt agtcataggg gaaggaatga ctgtgcttct gccattttgg 360
 ttttctgaag taaatctctg atatagttgc tgagtcagta gaatagtccc atcagccaca 420
 gattggattt ctataccaag aaaatattca aagtttccaa ttgtgtaaga caacaattgg 480
 aatgtatctt ggtggtgagt tgctgaatta ttacatg 517

<210> 1209
 <211> 366
 <212> DNA
 <213> Glycine max
 <223> unsure at all n locations
 <400> 1209

atgacgaaat ggctcacaag aggatgtaat gatctttcaa agtggagagg accttacaat 60
 atgtcaatac caaccacaga gagtccaatt aagacatcca tagtttcctt agtttcacaa 120
 tatcatatct taagttgggt atatataaat atcactatca tagcttacc acacagtgc 180
 aaagcataga catacactga ctctgtcaat ttattccac tatgatggga taaaaaactt 240
 ggcattatgt tcagttctac tatgtctcta ttgttctctt ctttctatgt gcttggttaat 300
 caatttttgc atgtactttc aactgtaata atangaaaat aattttcctt tatgggtctc 360
 ttctag 366

<210> 1210
 <211> 254
 <212> DNA
 <213> Glycine max
 <400> 1210

agcttaagag accacacctg atccttacac ttcactagct ggtcatagtg cttctgcac 60
 tcatgagtac ttcacacaaa actagagggg ctaccacagg ttgcttcaac accatgtatt 120
 ttgctgttat ctccaccact tatgatccta gtgatgaact ctggcatgac cttaatagca 180
 acccttccat cacaaccctt catgtactga atagggcgtg caccagactt gatacggggac 240
 ttgttagggc aaga 254